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(54) **COMPOSITIONS AND METHODS FOR COUNTERACTING FACTOR XA INHIBITION**

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(57) **ABSTRACT**

The disclosure provides compositions and methods for counteracting the effects of direct activated Factor X (FXa) inhibitors in a subject by administering a variant of FXa.

20 Claims, 17 Drawing Sheets

Specification includes a Sequence Listing.

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- Agenda of the MSTP Retreat held at Villanova University on Aug. 1, 2012 during which Nabil Thalji presented the slides in NPL Document #1 (above). Believed to correspond to document D3 in the ISR in PCT/IB2014/058494.
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- Non-Final Office action in U.S. Appl. No. 15/031,077 (dated Apr. 20, 2018).
- Amendment and Response to Non-Final Office Action in U.S. Appl. No. 15/031,077 (dated Oct. 22, 2018).
- Final Office Action in U.S. Appl. No. 15/031,077 (dated Jan. 11, 2019).
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* cited by examiner

FIG. 1A

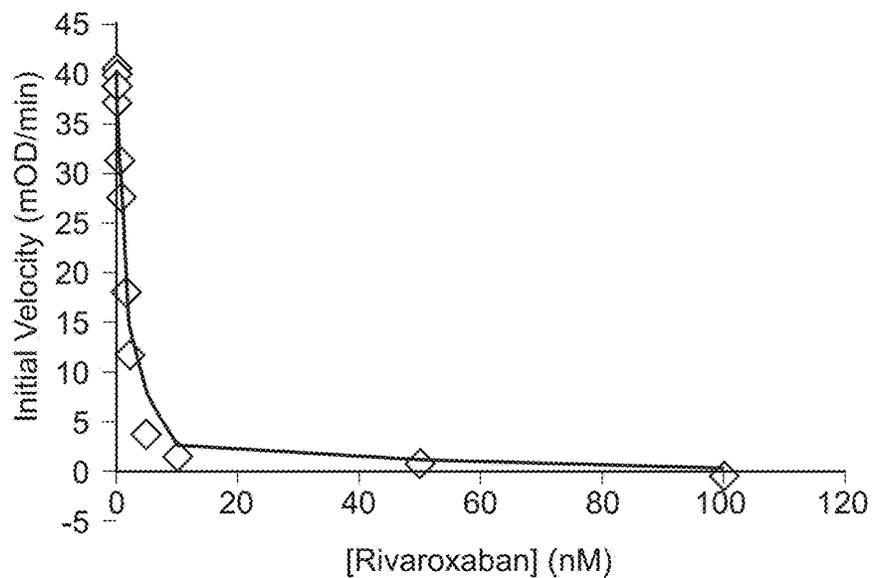


FIG. 1B

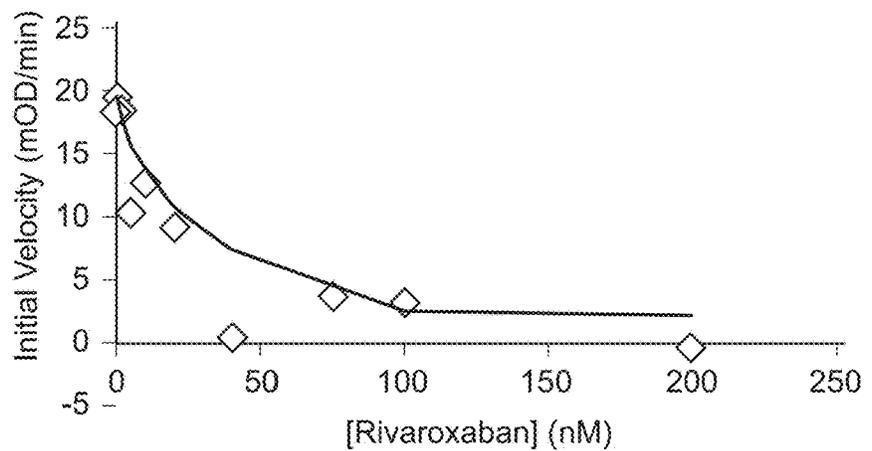


FIG. 2A

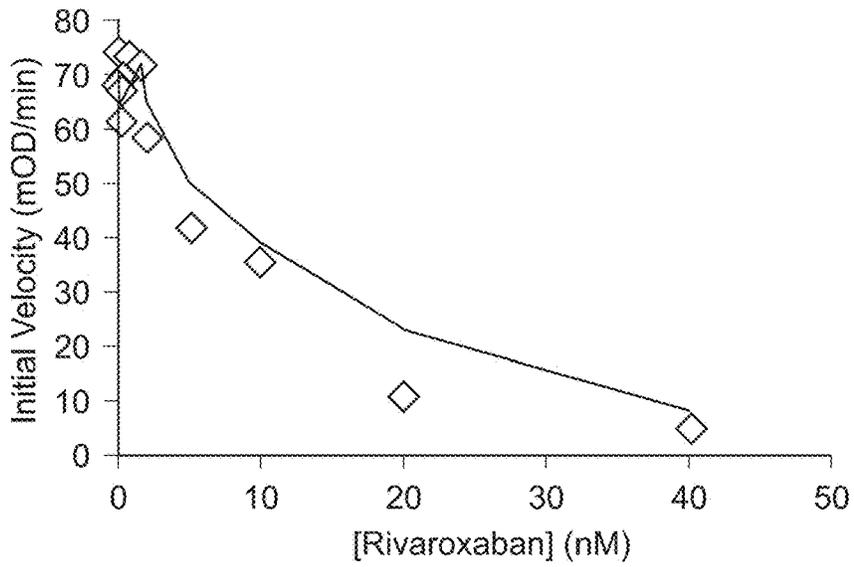


FIG. 2B

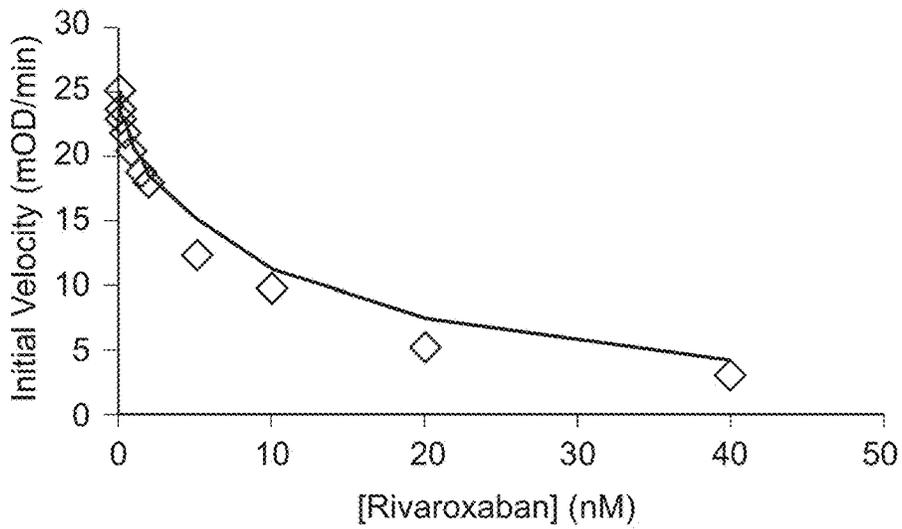


FIG. 3

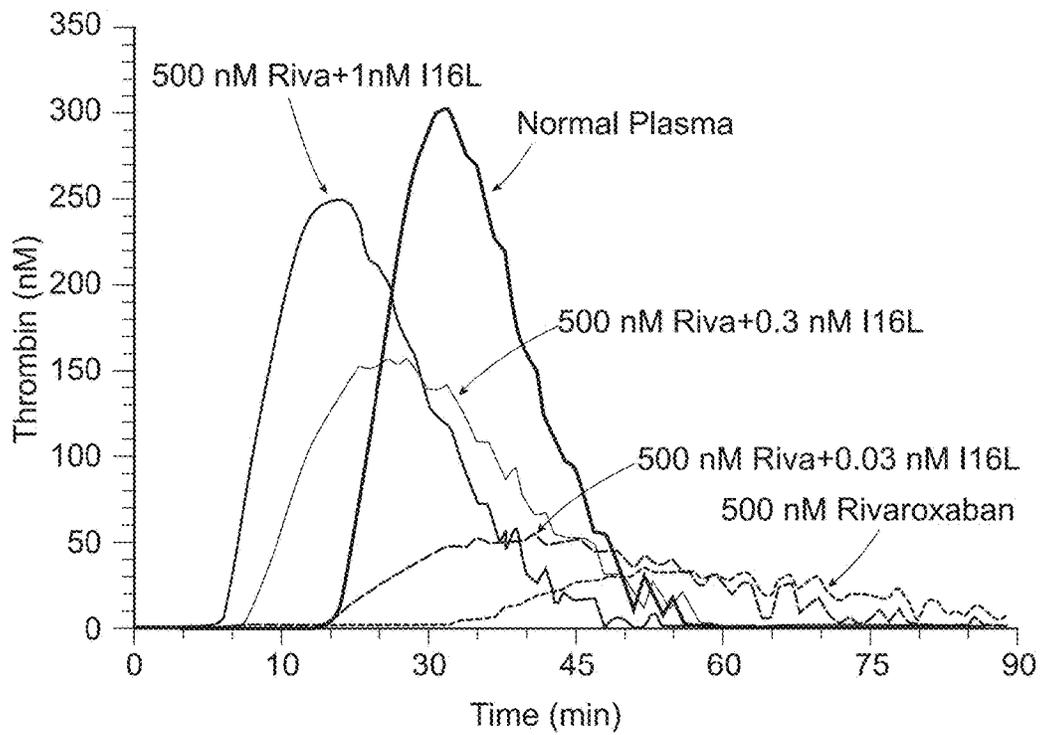


FIG. 4A

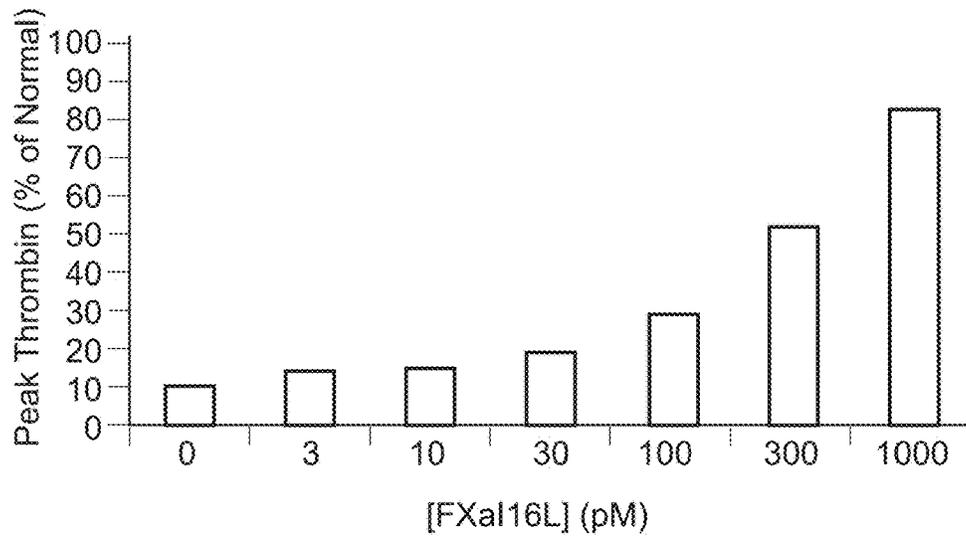


FIG. 4B

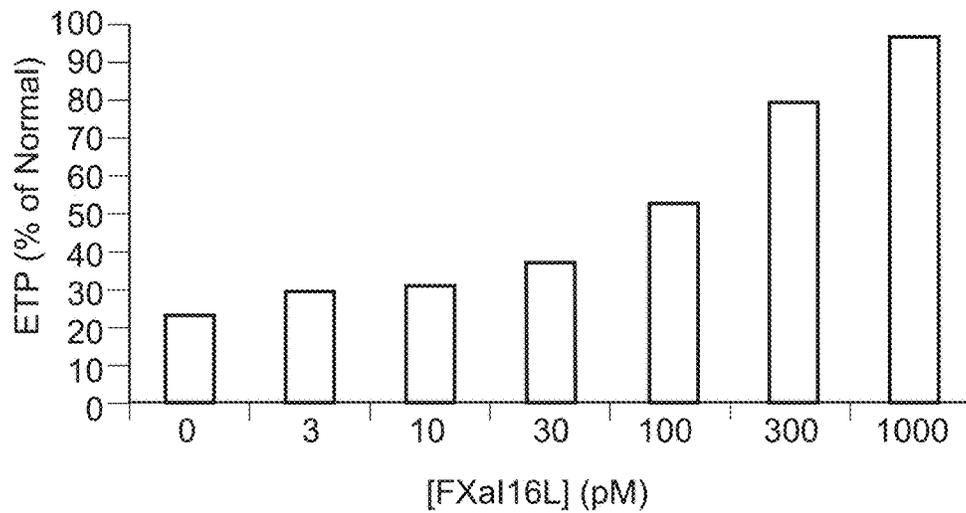


FIG. 4C

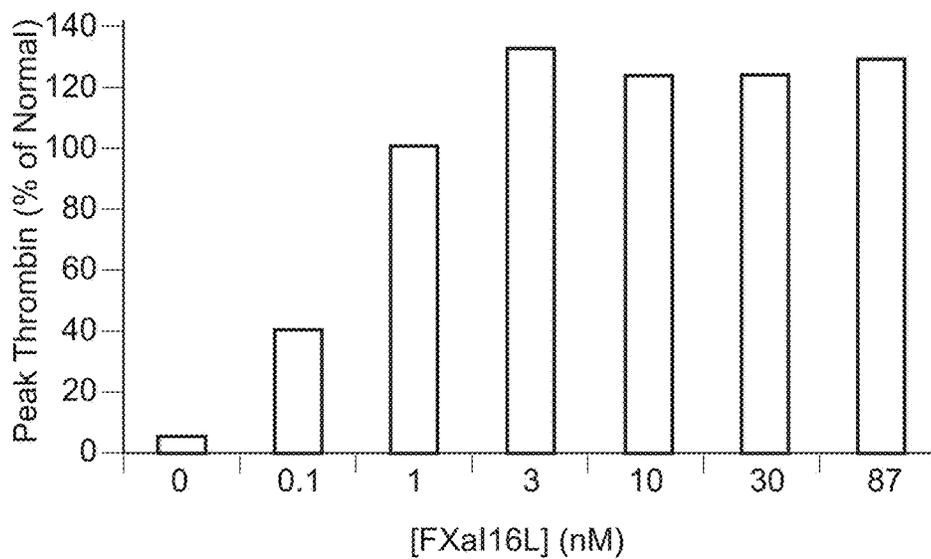


FIG. 4D

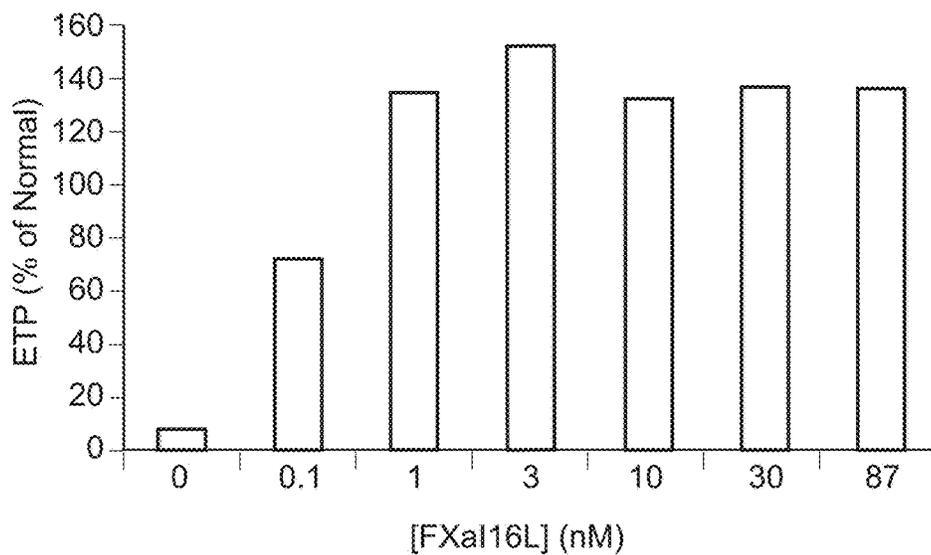


FIG. 5A

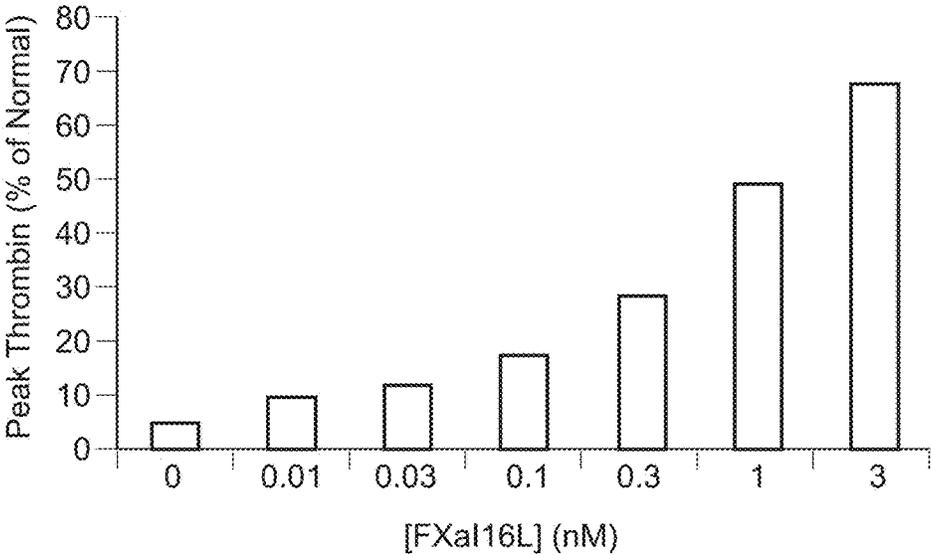


FIG. 5B

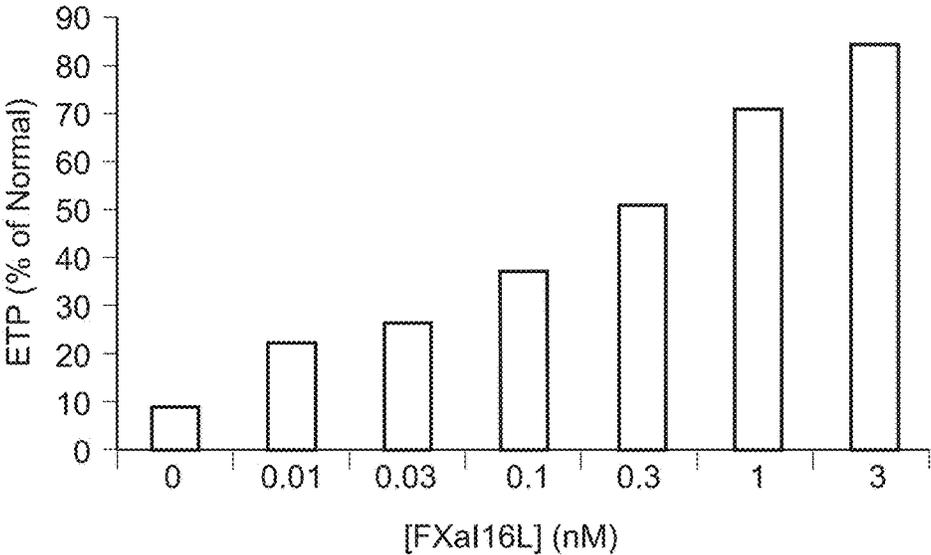


FIG. 6A

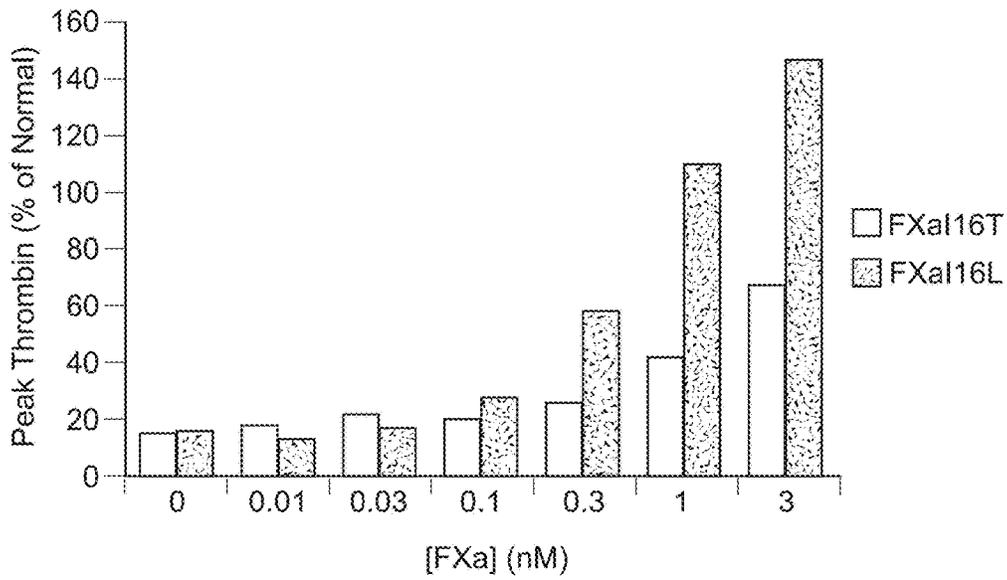


FIG. 6B

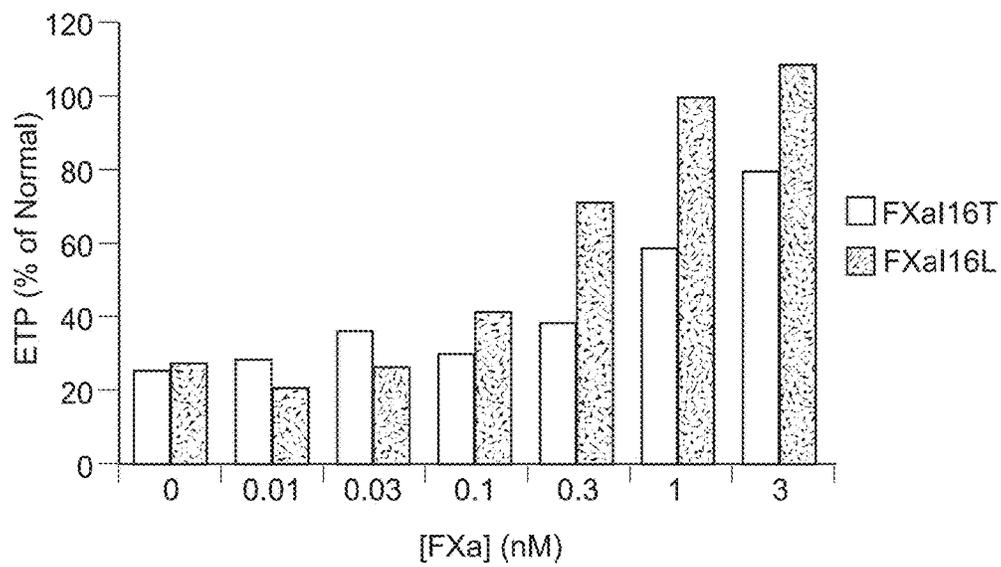


FIG. 7A

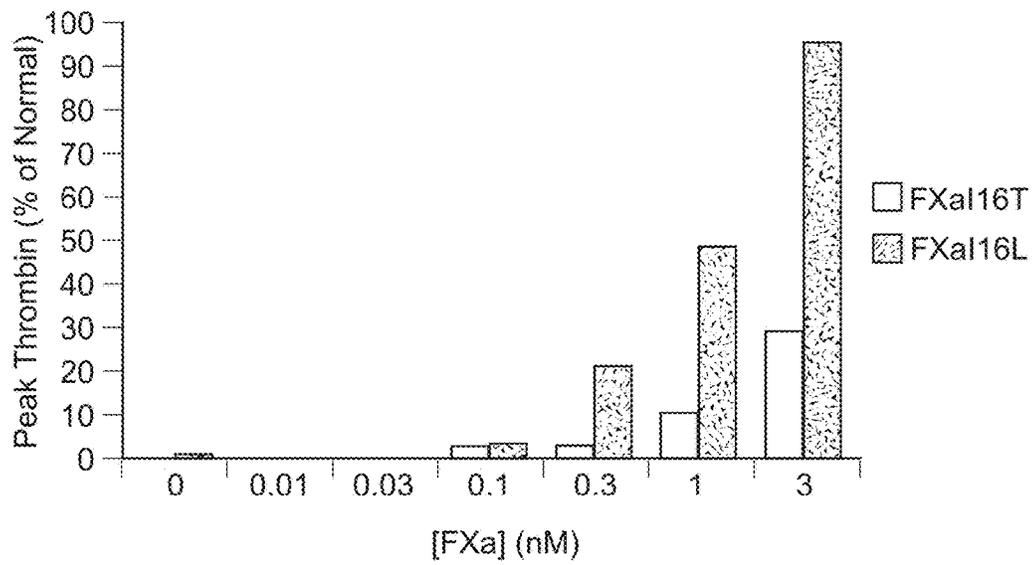
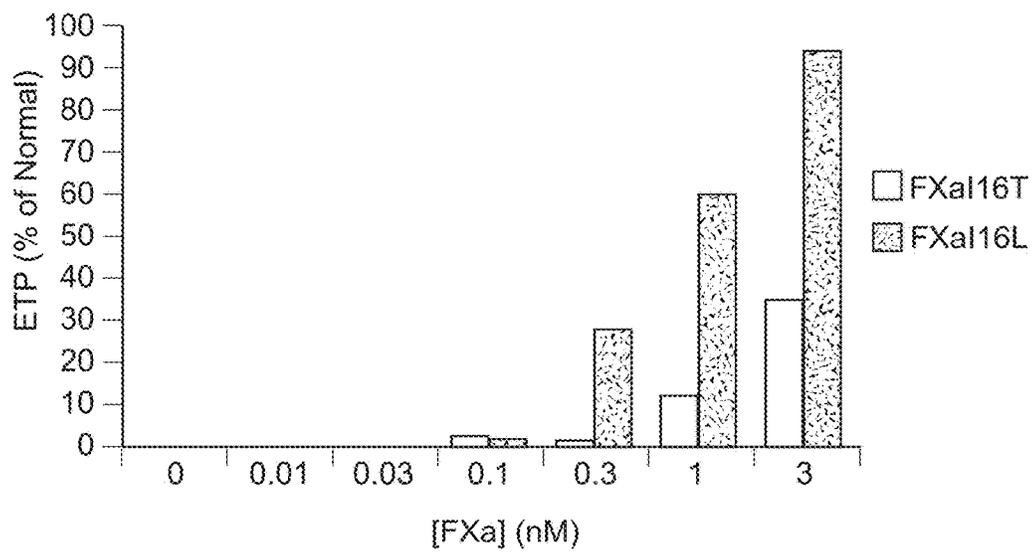


FIG. 7B



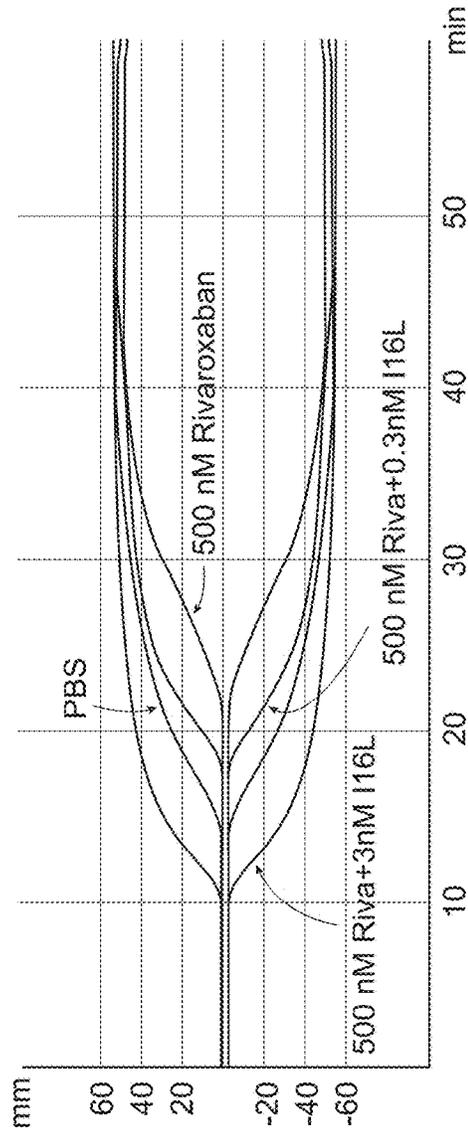


FIG. 8A

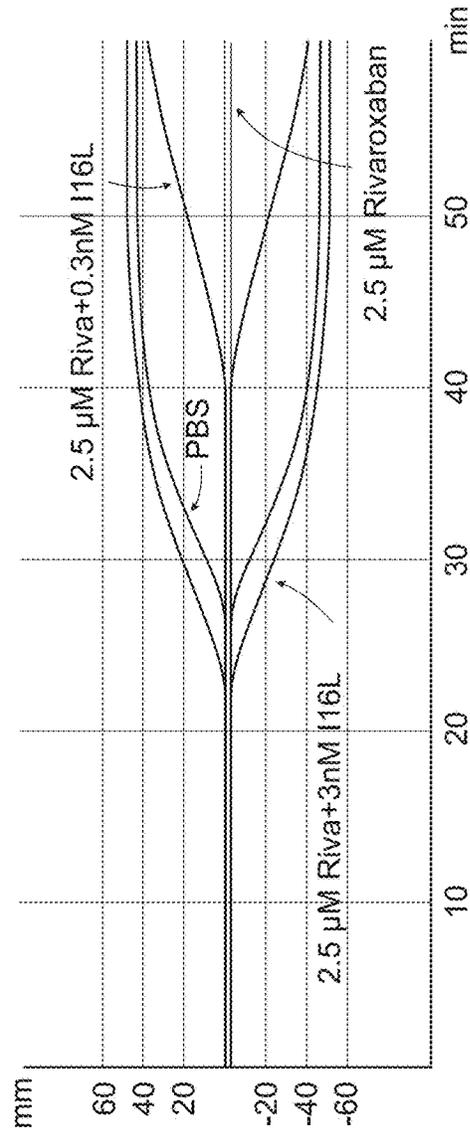


FIG. 8B

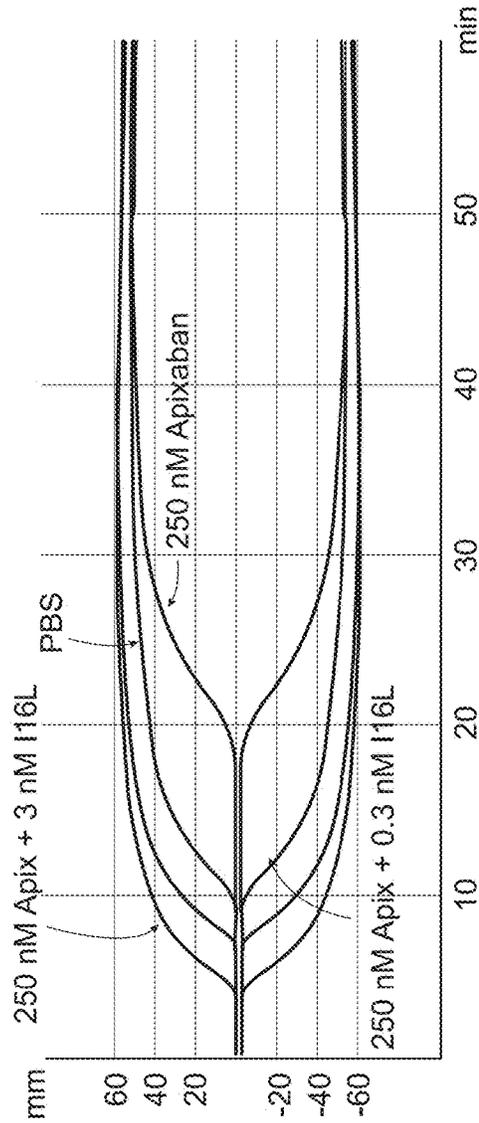


FIG. 9A

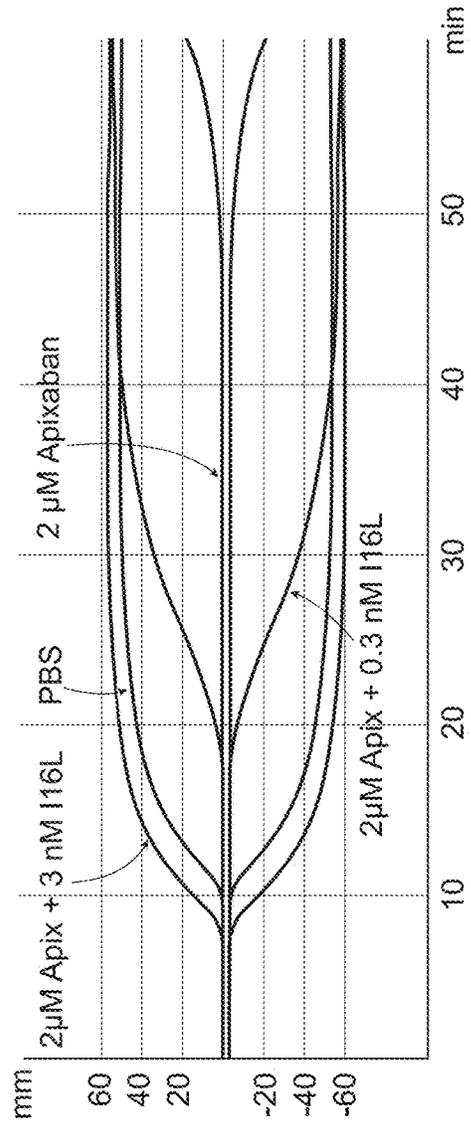


FIG. 9B

FIG. 10A

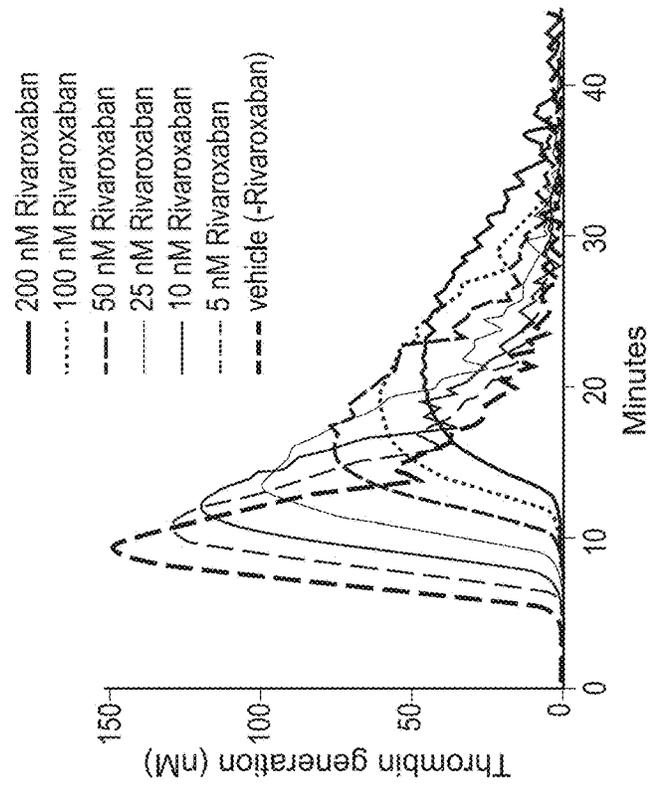


FIG. 10B

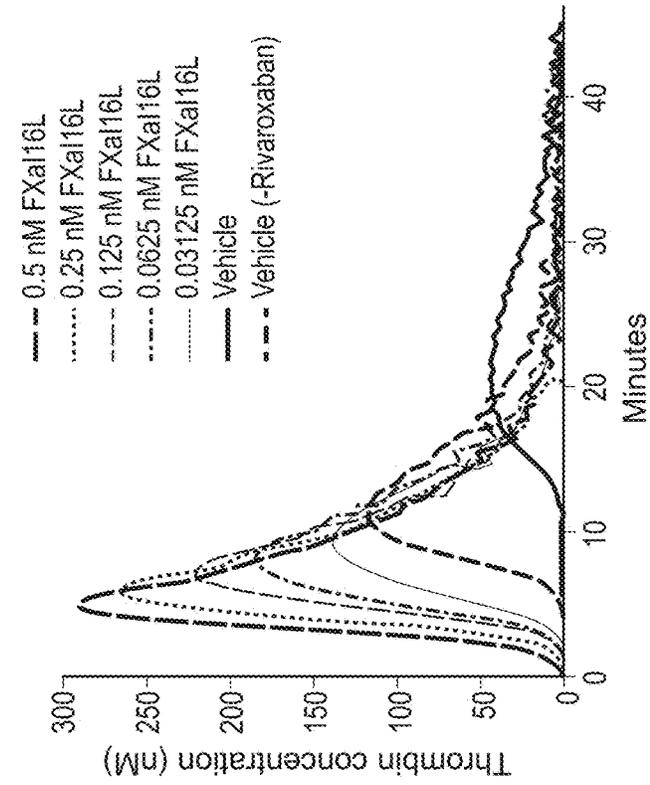


FIG. 11

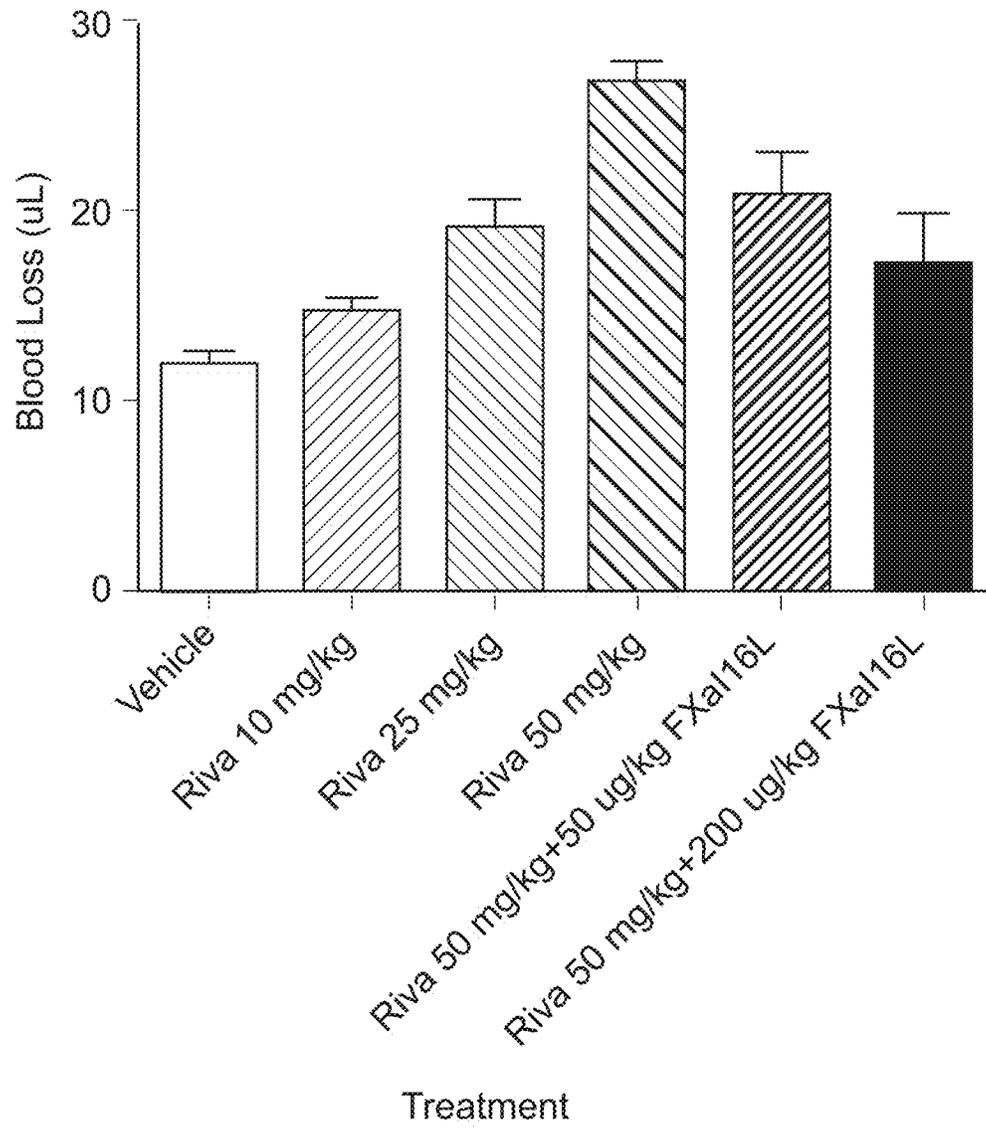


FIG. 12

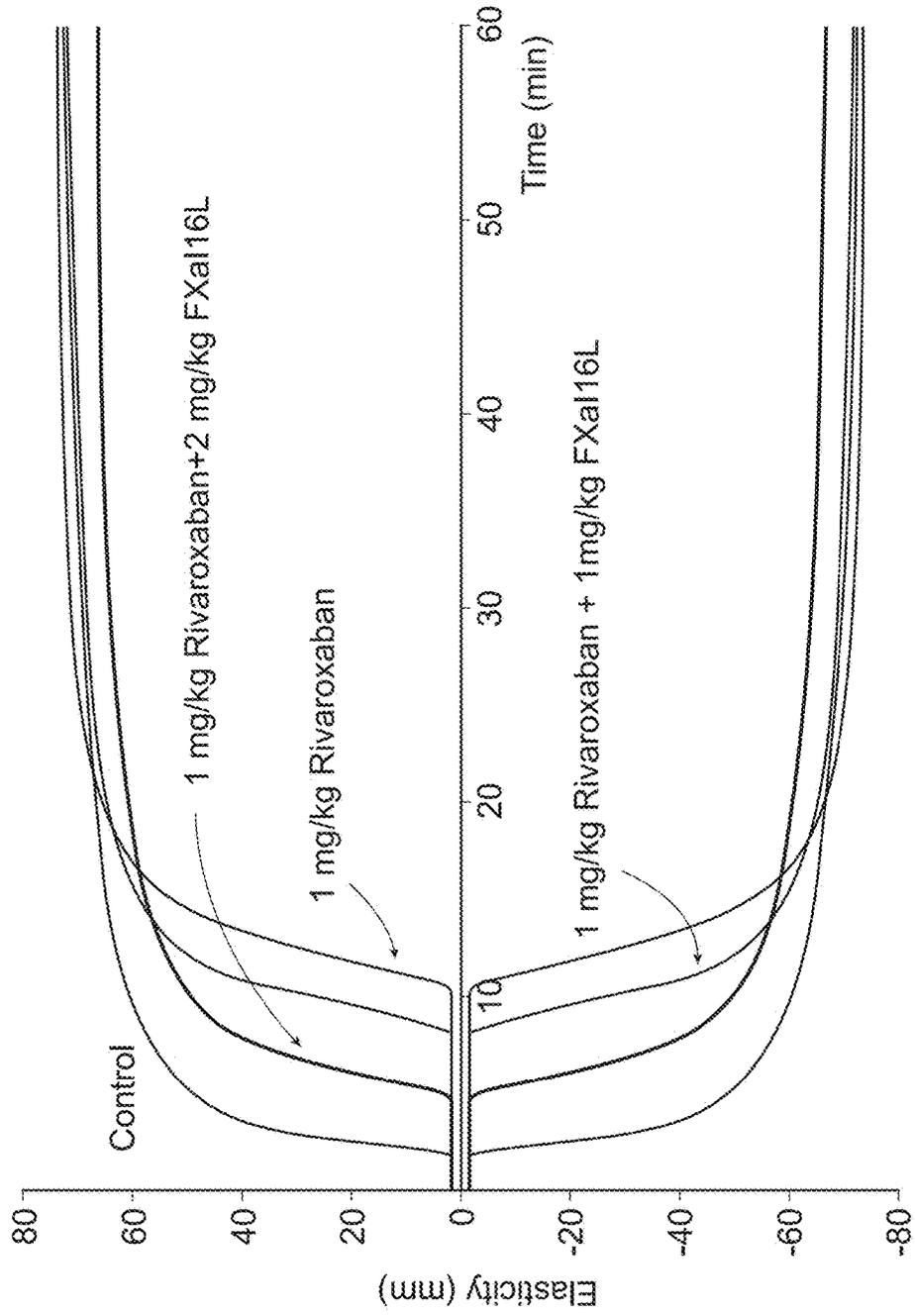


FIG. 13A

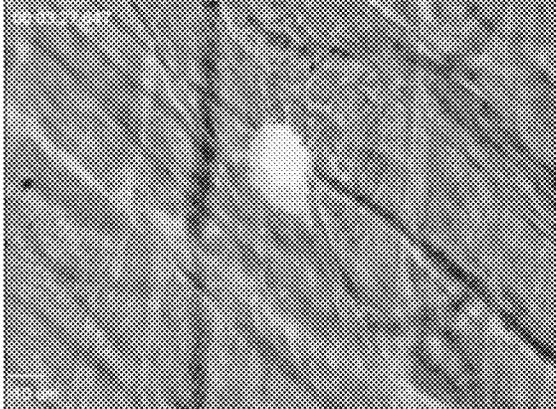


FIG. 13B

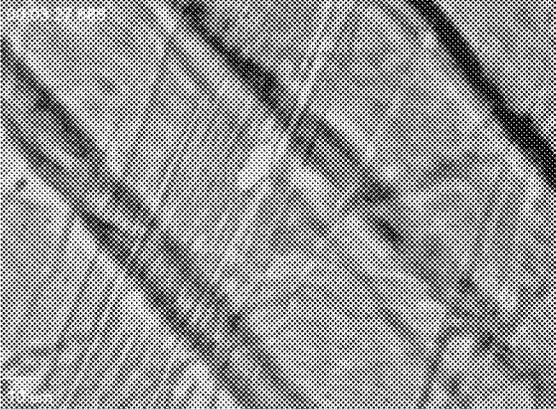


FIG. 13C

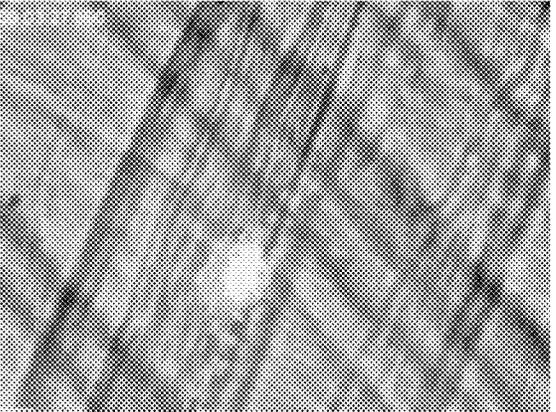


FIG. 14**Human Factor X Preprotein (SEQ ID NO:1)**

1 MGRPLHLVLL SASLAGLLL GESLFIRREQ ANNILARVTR ANSELEEMKK GHLEECMEE
61 TCSYEEAREV FEDSDKTNEF WNKYKDGQDC ETSPCQNOQK CKDGLGEYTC TCLEGFEGKN
121 CELFTRK LCS LDNGDCDQFC HEEQNSV VCS CARGYTLADN GKACIFTPGY PCGKQTLERR
181 KRSVAQATSS SGEAPDSITW KPYDAADLDP TENPFDLLDF NQTQPERGDN NLTRIVGGQE
241 CKDGECPWQA LLINEENEGF CGGTILSEFY ILTAAHCLYQ AKRFKVRVGD RNTEQEEGGE
301 AVHEVEVVIK HNRFTKETYD FDIAVLRLKT PITFRMNVAP ACLPERDWA E STLMTQKTGI
361 VSGFGRTHEK GRQSTRLKML EVPYVDRNSC KLSSSFIITQ NMFCAGYDTK QEDACQGD SG
421 GPHVTRFKDT YFVTGIVSWG EGCARKGKYG IYTKVTAFLK WIDRSMKTRG LPKAKSHAPE
481 VITSSPLK

FIG. 15**Human Factor X Preproteinein cDNA (SEQ ID NO:2)**

1 GACTTTGCTC CAGCAGCCTG TCCCAGTGAG GACAGGGACA CAGTACTCGG CCACACCATG
61 GGGCGCCAC TGCACCTCGT CFTGCTCAGT GCCTCCCTGG CTGGCCCTCT GCTGCTCGGG
121 GAAAGTCTGT TCATCCGCAG GGAGCAGGCC AACAAATCC TGGCGAGGGT CACGAGGGCC
181 AATTCCCTTC TTGAAGAGAT GAAGAAAGGA CACCTCGAAA GAGAGTGCAT GGAAGAGACC
241 TGCTCATACG AAGAGGCCCG CGAGGTCTTT GAGGACAGCG ACAAGACGAA TGAATTCTGG
301 AATAAATACA AAGATGGCGA CCAGTGTGAG ACCAGTCTTT GCCAGAACCA GGGCAAATGT
361 AAAGACGGCC TCGGGGAATA CACCTGCACC TGTTTAGAAG GATTCGAAGG CAAAAACTGT
421 GAATTATTCA CACGGAAGCT CTGCAGCCTG GACAAACGGG ACTGTGACCA GTTCTGCCAC
481 GAGGAACAGA ACTCTGTGGT GTGCTCCTGC GCCCGCGGGT ACACCCCTGGC TGACAACGGC
541 AAGGCCGTGA TTCCCACAGG GCCCTACCCC TGTGGGAAAC AGACCCTGGA ACGCAGGAAG
601 AGGTCAGTGG CCCAGGCCAC CAGCAGCAGC GGGGAGGCC CTGACAGCAT CACATGGAAG
661 CCATATGATG CAGCCGACCT GGACCCACC GAGAACCCCT TCGACCTGCT TGACTTCAAC
721 CAGACGCAGC CTGAGAGGGG CGACAACAAC CTCACCAGGA TCGTGGGAGG CCAGGAATGC
781 AAGGACGGG AGTGTCCCTG GCAGGCCCTG CTCATCAATG AGGAAAACGA GGGTTTCTGT

FIG. 15 continued

841 GGTGGAACCA TTCTGAGCGA GTTCTACATC CTAACGGCAG CCCACTGTCT CTACCAAGCC
901 AAGAGATTCA AGGTGAGGGT AGGGACCCGG AACACGGAGC AGGAGGAGGG CCGTGAGGGC
961 GTGCACGAGG TGGAGGTGGT CATCAAGCAC AACCGGTTCA CAAAGGAGAC CTATGACTTC
1021 GACATCGCCG TGCTCCGGCT CAAGACCCCC ATCACTTCC GCATGAACGT GGCGCCTGCC
1081 TGCCFCCCG AGCGTGACTG GGCCGAGTCC ACCGTGATGA CGCAGAAGAC GGGATTGTG
1141 AGCGGCTTCG GGCGCACCCA CGAGAAGGC CGGCAGTCCA CCAGGCTCAA GATGCTGGAG
1201 GTGCCCTACG TGGACCCGAA CAGCTGCAAG CTGTCCAGCA GCTTCATCAT CACCCAGAAC
1261 ATGTTCTGTG CCGGCTACGA CACCAAGCAG GAGGATGCC T GCCAGGGGGA CAGCGGGGGC
1321 CCGCACGTCA CCCGCTTCAA GGACACCTAC TTCGTGACAG GCATCGTCAG CTGGGAGAG
1381 GGCTGTGCC GTAAGGGAA GTACGGGATC TACACCAAGG TCACCGCCCTT CCTCAAGTGG
1441 ATCGACAGGT CCATGAAAAC CAGGGGCTTG CCCAAGGCCA AGAGCCAATGC CCCGGAGGTC
1501 ATAACGTCTT CTCCATTAAA GTGAGATCCC ACTCAAAAAA AAAAAAAA AAAAAAAA

1

COMPOSITIONS AND METHODS FOR COUNTERACTING FACTOR XA INHIBITION

CROSS-REFERENCE TO RELATED APPLICATION

This application claims the benefit of U.S. Provisional Application No. 61/759,332, filed 31 Jan. 2013, the contents of which are incorporated herein by reference in its entirety.

REFERENCE TO SEQUENCE LISTING

The Sequence Listing submitted concurrently herewith under 37 CFR § 1.821 in a computer readable form (CRF) via EFS-Web as file name PC72006_SEQLIST_ST25.txt is incorporated herein by reference. The electronic copy of the Sequence Listing was created on 17 Jan. 2018, with a file size of 6,835 bytes.

BACKGROUND OF THE INVENTION

Pharmacological anticoagulation is the mainstay of treatment for patients with prothrombotic conditions. For over fifty years, the only oral anticoagulant available was warfarin, an inhibitor of the vitamin K epoxide reductase (VKOR) that recycles oxidized vitamin K. Warfarin has many drawbacks, including unpredictable pharmacokinetics that necessitate frequent monitoring of coagulation parameters and dose adjustment. However, in the event of emergency bleeding or the need for urgent surgery, antidotes exist that allow rapid and complete reversal.

Oral direct FXa inhibitors are emerging anticoagulants that have the potential to simplify dosing schemes and hemostatic monitoring in patients with prothrombotic diseases when compared to standard treatments, such as warfarin. Although these drugs have many advantages over warfarin, no fully efficacious reversal agent is available for these novel anticoagulants.

The lack of a specific countermeasure to their effects, however, is a critical unmet clinical need that could limit the widespread adoption of these agents due to fears of unmanageable bleeding.

SUMMARY OF THE INVENTION

Applicants have addressed this critical unmet clinical need by providing compositions and methods for counteracting the effects of direct activated Factor X (FXa) inhibitors.

According to some embodiments, the disclosure provides methods for reducing or preventing bleeding in a subject being treated with a direct Factor Xa (FXa) inhibitor by administering a composition comprising a Factor Xa variant containing at least one modification including substitution for the wild-type amino acid at position 16 (using the chymotrypsin numbering system) with Thr, Leu, Phe, Asp or Gly, or substitution for the wild-type amino acid at position 17 (using the chymotrypsin numbering system) with Leu, Ala, or Gly. In certain embodiments, treatment with a composition comprising a FXa variant results in at least a 50% reduction in bleeding. According to certain embodiments, direct Factor Xa inhibitors include rivaroxaban or apixaban. In some embodiments, the plasma concentration of the direct FXa inhibitor is a typical therapeutic amount or a suprathreshold amount. For example, in some embodiments, the plasma concentration of rivaroxaban can be about

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500 nM, or greater, and the plasma concentration of apixaban can be about 250 nM, or greater. According to certain embodiments the FXa variant contains the substitution I16L. In some embodiments, the FXa variant is capable of countering the effect of the direct Factor Xa inhibitor at a plasma concentration that is at least 100-fold lower than the plasma concentration of the Factor Xa inhibitor. In certain embodiments, the composition comprising the FXa variant is administered before a planned surgery, after an injury, or after an intentional or accidental overdose with a direct FXa inhibitor. In some embodiments, hemostasis in the subject is monitored using a hemostasis assay after a first dose with a FXa variant and, if adequate hemostasis is not attained by a predetermined time, at least one second dose of FXa variant is administered to achieve sufficient hemostasis. According to some embodiments, the predetermined time is about 15 mins, 30 mins, 45 mins or 60 mins after the first dose of FXa variant is administered. Other times are also possible. In some other embodiments, at least a second procoagulant is administered in addition to FXa variant, including for example, a different FXa variant, factor IX, factor XIa, factor XIIa, factor VIII, factor VIIa, FEIBA or prothrombin complex concentrate (PCC).

According to some embodiments, the disclosure provides methods for increasing the amount of thrombin produced in response to activation of the extrinsic or intrinsic clotting pathway in a subject being treated with a direct Factor Xa (FXa) inhibitor by administering a composition comprising a Factor Xa variant containing at least one modification including substitution for the wild-type amino acid at position 16 (using the chymotrypsin numbering system) with Thr, Leu, Phe, Asp- or Gly, or substitution for the wild-type amino acid at position 17 (using the chymotrypsin numbering system) with Leu, Ala, or Gly. According to certain embodiments, direct Factor Xa inhibitors include rivaroxaban or apixaban. In some embodiments, the plasma concentration of the direct FXa inhibitor is a typical therapeutic amount or a suprathreshold amount. For example, in some embodiments, the plasma concentration of rivaroxaban can be about 500 nM, or greater, and the plasma concentration of apixaban can be about 250 nM, or greater. According to certain embodiments the FXa variant contains the substitution I16L. According to certain embodiments, the amount of thrombin produced increases by about 5%, 10%, 15%, 20%, 30%, 40%, 50%, 60%, 70%, 80%, 90%, 100%, 150%, 200%, or more. In some embodiments, the FXa variant is capable of countering the effect of the direct Factor Xa inhibitor at a plasma concentration that is at least 100-fold lower than the plasma concentration of the Factor Xa inhibitor. In certain embodiments, the composition comprising the FXa variant is administered before a planned surgery, after an injury, or after an intentional or accidental overdose with a direct FXa inhibitor. In some embodiments, hemostasis in the subject is monitored using a hemostasis assay after a first dose with a FXa variant and, if adequate hemostasis is not attained by a predetermined time, at least one second dose of FXa variant is administered to achieve sufficient hemostasis. According to some embodiments, the predetermined time is about 15 mins, 30 mins, 45 mins or 60 mins after the first dose of FXa variant is administered. Other times are also possible. In some other embodiments, at least a second procoagulant is administered in addition to FXa variant, including for example, a different FXa variant, factor IX, factor XIa, factor XIIa, factor VIII, factor VIIa, FEIBA or prothrombin complex concentrate (PCC).

According to some embodiments, the disclosure provides methods for decreasing clotting time (as measured, for

example, using PT or INR, or some other assay) in a subject being treated with a direct Factor Xa (FXa) inhibitor by administering a composition comprising a Factor Xa variant containing at least one modification including substitution for the wild-type amino acid at position 16 (using the chymotrypsin numbering system) with Thr, Leu, Phe, Asp or Gly, or substitution for the wild-type amino acid at position 17 (using the chymotrypsin numbering system) with Leu, Ala, or Gly. According to certain embodiments, direct Factor Xa inhibitors include rivaroxaban or apixaban. In some embodiments, the plasma concentration of the direct FXa inhibitor is a typical therapeutic amount or a suprathereapeutic amount. For example, in some embodiments, the plasma concentration of rivaroxaban can be about 500 nM, or greater, and the plasma concentration of apixaban can be about 250 nM, or greater. According to certain embodiments the FXa variant contains the substitution I16L. According to certain embodiments, clotting time is reduced by about 5%, 10%, 15%, 20%, 30%, 40%, 50%, 60%, 70%, 80%, 90%, or more. In some embodiments, the FXa variant is capable of countering the effect of the direct Factor Xa inhibitor at a plasma concentration that is at least 100-fold lower than the plasma concentration of the Factor Xa inhibitor. In certain embodiments, the composition comprising the FXa variant is administered before a planned surgery, after an injury, or after an intentional or accidental overdose with a direct FXa inhibitor. In some embodiments, hemostasis in the subject is monitored using a hemostasis assay after a first dose with a FXa variant and, if adequate hemostasis is not attained by a predetermined time, at least one second dose of FXa variant is administered to achieve sufficient hemostasis. According to some embodiments, the predetermined time is about 15 mins, 30 mins, 45 mins or 60 mins after the first dose of FXa variant is administered. Other times are also possible. In some other embodiments, at least a second procoagulant is administered in addition to FXa variant, including for example, a different FXa variant, factor IX, factor XIa, factor XIIa, factor VIII, factor Vila, FEIBA or prothrombin complex concentrate (PCC).

BRIEF DESCRIPTION OF THE DRAWINGS

FIGS. 1A-B show inhibition of free wt-FXa or FXa^{I16L} by rivaroxaban. The initial velocity of peptidyl substrate (SpecXa; 200 uM) hydrolysis by A) wt-FXa (2 nM) or B) FXa^{I16L} (6 nM) was determined at increasing concentrations of rivaroxaban. The Ki value is given on each graph.

FIGS. 2A-B show rivaroxaban inhibition of wt-FXa or FXa^{I16L} assembled in prothrombinase. The initial velocity of peptidyl substrate (SpecXa; 200 uM) hydrolysis by A) wt-FXa (2 nM) or B) FXa^{I16L} (6 nM) in the presence of PCPS (20 uM) and FVa (40 nM) was determined at increasing concentrations of rivaroxaban.

FIG. 3 shows the effect of different concentrations of FXa^{I16L} on reversing the effects on thrombin generation of rivaroxaban.

FIGS. 4A-D show the effect of FXa^{I16L} on reversing the effects of rivaroxaban. Normal human plasma was incubated with 500 nM rivaroxaban and in the absence or presence of increasing concentrations of FXa^{I16L}. Following data analysis, peak thrombin (A and C) and total thrombin generated (ETP; B and D) were plotted.

FIGS. 5A-B show FXa^{I16L} reverses the effects of high dose rivaroxaban. Normal human plasma was incubated with 7.5 uM rivaroxaban and in the absence or presence of

increasing concentrations of FXa^{I16L}. Following data analysis, peak thrombin (A) and total thrombin generated (ETP; B) were plotted.

FIGS. 6A-B show FXa^{I16L} or FXa^{I16T} reverses the effects of 250 nM apixaban. Normal human plasma was incubated with 250 nM apixaban and in the absence or presence of increasing concentrations of FXa^{I16L} or FXa^{I16T}. Following data analysis, peak thrombin (A) and total thrombin generated (ETP; B) were plotted.

FIGS. 7A-B show FXa^{I16L} or FXa^{I16T} reverses the effects of high dose apixaban. Normal human plasma was incubated with 2.0 uM Apixaban and in the absence or presence of increasing concentrations of FXa^{I16L} or FXa^{I16T}. Following data analysis, peak thrombin (A) and total thrombin generated (ETP; B) were plotted.

FIGS. 8A-B show FXa^{I16L} corrects whole blood clotting in the presence of rivaroxaban. Whole blood thromboelastography was used to assess the ability of FXa^{I16L} to reverse the effects of rivaroxaban at a typical (A) and a high (B) dose.

FIGS. 9A-B show FXa^{I16L} corrects whole blood clotting in the presence of apixaban. Whole blood thromboelastography was used to assess the ability of FXa^{I16L} to reverse the effects of apixaban at a typical (A) and a high (B) dose.

FIGS. 10A-B show that FXa^{I16L} counteracts rivaroxaban in a thrombin generation assay. FIG. 10A shows a dose response of rivaroxaban and FIG. 10B shows a dose response of FXa^{I16L} in the presence of rivaroxaban.

FIG. 11 shows that FXa^{I16L} counteracts rivaroxaban in a mouse tail clip bleeding model.

FIG. 12 demonstrates that rivaroxaban administered to mice delays clotting time of whole blood measured using ROTEM and that administration with FXa^{I16L} dose-responsively counteracts the effect of rivaroxaban.

FIGS. 13A, 13B, and 13C show that rivaroxaban administered to a mouse prevents clot formation at a site of vascular injury in the cremaster muscle caused by laser and that administration with FXa^{I16L} counteracts the effect of rivaroxaban. Clot formation was visualized using intravital microscopy and fluorescently labeled antibodies against fibrin and platelets. FIG. 13A shows clot formation in an untreated mouse. FIG. 13B shows that rivaroxaban delayed and reduced platelet accumulation and prevented fibrin deposition. By contrast, FIG. 13C shows that in a mouse administered rivaroxaban and FXa^{I16L} clot formation occurred at the site of injury.

FIG. 14 is the amino acid sequence of wild-type human Factor X preprotein (SEQ ID NO:1). The signal peptide corresponds to amino acids 1-23. The propeptide corresponds to amino acids 24-40. The mature light chain of activated Factor X (FXa) corresponds to amino acids 41-179. The mature heavy chain of activated FXa (after removal of the activation peptide) corresponds to amino acids 235-488. The activation peptide (AP) corresponds to amino acids 183-234.

FIG. 15 is the nucleotide sequence of the cDNA encoding wild-type human Factor X preprotein (SEQ ID NO:2). The coding sequence corresponds to nucleotides 58 to 1524.

DETAILED DESCRIPTION

The disclosure provides compositions and methods for counteracting the anti-coagulant effect of a direct FXa inhibitor in a subject in need thereof. Applicants have discovered that certain FXa variants rapidly and completely counteract the effect of a direct FXa inhibitor in a dose dependent manner. More specifically, applicants have dis-

covered that a relatively small amount of an FXa variant restores normal coagulation activity in vivo in the presence of FXa inhibitor at therapeutic concentrations and even at supratherapeutic concentrations. By providing fast and effective antidotes for the anti-coagulation effects of direct FXa inhibitors, Applicants' disclosure therefore contributes to fulfilling the promise of these advantageous anti-coagulants.

Coagulation factor X (FX) is a zymogen which, upon activation by the intrinsic factor IX/factor VIII or extrinsic pathway (tissue factor/factor VIIa), becomes FXa, which is the protease moiety of prothrombinase. Following proteolytic cleavage of the Arg-Ile scissile bond, releasing an activation peptide (AP), a series of well defined structural changes in the zymogen drives the activation process to the mature active serine protease (See Toso et al., (2008) *J. Biol. Chem.* 283, 18627-18635; Bunce et al., (2011) *Blood* 117, 290-298; and Ivanciu et al., (2011) *Nat. Biotechnol.* 29, 1028-1033, incorporated by reference herein in their entirety). The mature FXa has a light chain and a heavy chain that contains the catalytic domain. The mature FXa becomes an active serine protease upon formation of the prothrombinase complex, which includes binding of an activated cofactor, Factor Va (FVa).

Variant forms of FX have been developed that upon activation cleavage yield a zymogen-like FXa variant. That is, once cleaved, the resulting FXa variant has poor active site function and is more resistant to inactivation by circulating inhibitors (i.e. antithrombin III and TFPI). The FXa variants, thus, have longer half-lives in plasma than wild-type FXa. The FXa variant binds FVa, lipid membrane and calcium to form a fully active prothrombinase complex that efficiently activates prothrombin.

The zymogen-like variants of FXa circulate in the zymogen-like conformation and do not seem to be thrombogenic (See Toso et al., (2008) *J. Biol. Chem.* 283, 18627-18635 and Ivanciu et al., (2011) *Nat. Biotechnol.* 29, 1028-1033, incorporated by reference herein in their entirety). Examples of such FXa variants are described in International patent publication WO2007/059513, incorporated herein by reference in its entirety.

The enzymes of coagulation are trypsin-like enzymes that belong to the S1 peptidase family of proteases that bear a chymotrypsin-like fold. The coagulation proteases contain catalytic domains that are highly homologous to each other and to the ancestral serine proteases of digestion. The structural homology/identity is so great (>70%) that residues in the catalytic domains of the coagulation enzymes (including Factor Xa) are numbered according to the corresponding residues in chymotrypsinogen. (Chymotrypsin numbering system; see Bajaj and Birktoft, *Methods Enzymol.* 1993; 222:96-128, Table 2, and Bode W, Mayr I, Bauman Y, et al. The refined 1.9 Å crystal structure of human alpha-thrombin: interaction with D-Phe-Pro-Arg chloromethylketone and significance of the Tyr-Pro-Trp insertion segment. *EMBO J* 1989; 8(11):3467-3475, both of which are incorporated by reference herein in their entirety). Accordingly, amino acids may be referred to herein according to the chymotrypsin numbering system, which is well-known to those of skill in the art.

According to the disclosure, an FXa variant may be an FXa protein comprising an amino acid substitution that makes the variant more zymogen-like compared to a wild-type FXa protein in vivo or in vitro. FXa variants of the disclosure substantially regain wild-type FXa activity upon formation of prothrombinase. Examples of FXa variants that are useful in methods of the disclosure are variants com-

prising a modification selected from the group consisting of: a) Ile at position 16 is Thr, Leu, Phe, Asp or Gly and b) Val at position 17 is Leu, Ala, or Gly, according to the chymotrypsin numbering system. Amino acids 16 and 17 in the chymotrypsin numbering system occur at amino acids 235 and 236, respectively, of SEQ ID NO:1 (human Factor X preproprotein). In certain embodiments, FXa variants are FXa^{N16L} and FXa^{N16T} (the nomenclature used herein for the FXa variants recites the original amino acid at the numbered position according to the chymotrypsin numbering system followed by the substituted amino acid). The FXa variants can be variants of any mammalian FXa. Of particular interest, however, are FXa variants of human FXa.

In certain embodiments, the FX variant that is activated into a variant FXa of the disclosure may be further modified by inserting a non-native intracellular proteolytic cleavage site. In a non-limiting example, to express "activated" zymogen-like FXa variants in mammalian cells, a non-native intracellular proteolytic cleavage site can be inserted between the Arg at position 234 of SEQ ID NO:1 (position 15 in the chymotrypsin numbering system) and the amino acid at the position corresponding to position 235 of SEQ ID NO:1 (position 16 in the chymotrypsin numbering system) in the variant FX zymogen. In certain embodiments, the non-native intracellular protease cleavage site is Arg-Lys-Arg or Arg-Lys-Arg-Arg-Lys-Arg (SEQ ID NO:3). These cleavage sites are efficiently recognized by proteases (PACE/furin-like enzymes) within the cell and are removed. This cleavage may result in a processed variant FXa in which the mature heavy chain of the molecule now begins at the amino acid at the position corresponding to position 235 of SEQ ID NO:1 (position 16 in the chymotrypsin numbering system). Introduction of this cleavage site at said position allows for the intracellular conversion of FX to FXa.

In certain embodiments the entire amino acid sequence of the FX variant activation peptide (AP) (i.e., amino acids 183-234 of SEQ ID NO:1) is replaced with a non-native intracellular protease cleavage site. According to certain embodiments, the non-native intracellular protease cleavage site is Arg-Lys-Arg or Arg-Lys-Arg-Arg-Lys-Arg (SEQ ID NO:3). As explained above, this modification allows for intracellular cleavage of the FX variant expressed by cells. The intracellular cleavage converts FX variant to activated zymogen-like FXa variant which is then secreted by cells for subsequent purification. This approach obviates the need for extracellular cleavage that would otherwise be required to activate the variant clotting factor, for example, after isolating the protein or just before blood clotting.

In certain embodiments, FXa variants of the disclosure are derived from FX variant preproteins comprising native wild-type human signal sequence and/or propeptide sequence. In other embodiments, FX signal sequences and/or propeptide from non-human species can be used in place of the corresponding native amino acid sequences. And in yet other embodiments, signal sequence and/or propeptide sequence from other human or non-human secreted proteins can be used in place of the corresponding native amino acid sequences.

In an exemplary embodiment, a FXa variant comprises amino acids 41-179 and amino acids 235-488 of SEQ ID NO:1, wherein the amino acid at position 235 (isoleucine in the wild-type sequence) is substituted with a different amino acid selected from the group consisting of threonine (Thr), leucine (Leu), phenylalanine (Phe), aspartic acid (Asp), or glycine (Gly). These substitutions can respectively be written using the nomenclature I235T, I235L, I235F, I235D and

I235G, where the first letter is the single letter code for isoleucine and the last letter is the single letter code for the amino acid being substituted into the wild-type sequence. Because position 235 of SEQ ID NO:1 is equivalent to position 16 in the chymotrypsin numbering system, the same substitutions can be written I16T, I16L, I16F, I16D and I16G. In an embodiment, a FXa variant comprises amino acids 41-179 and amino acids 235-488 of SEQ ID NO:1, wherein the amino acid at position 235 is substituted with Thr (i.e., I235T or I16T). In an embodiment, a FXa variant comprises amino acids 41-179 and amino acids 235-488 of SEQ ID NO:1, wherein the amino acid at position 235 is substituted with Leu (i.e., I235L or I16L). In an embodiment, a FXa variant comprises amino acids 41-179 and amino acids 235-488 of SEQ ID NO:1, wherein the amino acid at position 235 is substituted with Asp (i.e., I235D or I16D). In an embodiment, a FXa variant comprises amino acids 41-179 and amino acids 235-488 of SEQ ID NO:1, wherein the amino acid at position 235 is substituted with Gly (i.e., I235G or I16G).

According to another exemplary embodiment, a FXa variant comprises amino acids 41-179 and amino acids 235-488 of SEQ ID NO:1, wherein the amino acid at position 236 (valine in the wild-type sequence) is substituted with a different amino acid selected from the group consisting of leucine (Leu), alanine (Ala), or glycine (Gly). These substitutions can respectively be written using the nomenclature V236L, V236A, and V236G, where the first letter is the single letter code for valine and the last letter is the single letter code for the amino acid being substituted into the wild-type sequence. Because position 236 of SEQ ID NO:1 is equivalent to position 17 in the chymotrypsin numbering system, the same substitutions can be written V17L, V17A, and V17G. In an embodiment, a FXa variant comprises amino acids 41-179 and amino acids 235-488 of SEQ ID NO:1, wherein the amino acid at position 236 is substituted with Leu (i.e., V236L or V17L). In an embodiment, a FXa variant comprises amino acids 41-179 and amino acids 235-488 of SEQ ID NO:1, wherein the amino acid at position 236 is substituted with Ala (i.e., V236A or V17A). In an embodiment, a FXa variant comprises amino acids 41-179 and amino acids 235-488 of SEQ ID NO:1, wherein the amino acid at position 236 is substituted with Gly (i.e., V236G or V17G).

In other embodiments, FXa variants of the disclosure, including those specific variants described in the preceding paragraphs, can include various isoforms of the light and/or mature heavy chain of the protein. Non-limiting exemplary isoforms of the FXa variant mature heavy chain include the alpha and beta versions of the mature heavy chain. Jesty et al., J Biol Chem. 1975 Jun. 25; 250(12):4497-504, incorporated by reference herein. Compositions of the disclosure can include FXa variant proteins in which one or the other or both alpha and beta mature heavy chain isoforms are represented.

According to yet other embodiments, isoforms of FXa variant proteins, including those specific variants described in the preceding paragraphs, can include isoforms in which a variable number of amino acids (for example, 1, 2, 3, 4, 5, 6, or more amino acids) are either missing from or added to the carboxy terminus of the light chain and/or mature heavy chains of the protein.

According to certain embodiments, FXa variants of the disclosure include proteins with a certain minimal degree of

homology or sequence identity compared to the amino acid sequence of wild-type FXa in SEQ ID NO:1. Thus, for example, FXa variants include proteins that contain a light and mature heavy chain that are at least 60%, 70%, 80%, 85%, 90%, 95%, 98%, or 99% homologous or identical in sequence with the wild-type FXa light and mature heavy chains in SEQ ID NO:1, wherein such FXa variants also include a substitution at the amino acid position corresponding to position 235 of SEQ ID NO:1 with Thr, Leu, Phe, Asp, or Gly, or a substitution at the amino acid position corresponding to position 236 of SEQ ID NO:1 with Leu, Ala, or Gly, and further wherein such FXa variants are zymogenic until incorporated into prothrombinase complex. In the amino acid sequence of SEQ ID NO:1, the wild-type FXa light chain sequence corresponds to amino acids 41 to 179 and the wild-type FXa mature heavy chain sequence corresponds to amino acids 235 to 488. Percentage amino acid sequence homology or identity can readily be determined using software such as Protein BLAST available at the website of the National Center for Biotechnology Information (<http://blast.ncbi.nlm.nih.gov/Blast.cgi>).

According to other non-limiting embodiments, FXa variants of the disclosure can also include FXa variants containing one or more post-translational modifications including, without limitation, one or more O-linked or N-linked carbohydrate groups or a variable number of gamma-carboxyglutamic acid (Gla) residues. FXa variants of the disclosure can further include chemically modified FXa variant proteins. Other FXa variants useful in the methods of the disclosure are also possible.

As used herein, the term FXa^{I16x} refers to a variant of activated Factor X wherein the amino acid corresponding to position 235 in SEQ ID NO:1 (corresponding to position 16 in the chymotrypsin numbering system) is changed from the amino acid in the wild-type sequence (isoleucine) to a different amino acid denoted "x". In some non-limiting exemplary embodiments, amino acid "x" can be threonine (Thr or T), leucine (Leu or L), phenylalanine (Phe or F), aspartic acid (Asp or D), or glycine (Gly or G).

As used herein, the term FXa^{V17y} refers to a variant of activated Factor X wherein the amino acid corresponding to position 236 in SEQ ID NO:1 (corresponding to position 17 in the chymotrypsin numbering system) is changed from the amino acid in the wild-type sequence (valine) to a different amino acid denoted "y". In some non-limiting exemplary embodiments, amino acid "y" can be leucine (Leu or L), alanine (Ala or A), or glycine (Gly or G).

The terms FXa^{I16x} and FXa^{V17y} are not limited by the protein sequence set forth in SEQ ID NO:1. Rather these terms additionally include the variety of isoforms and homologous proteins described herein with the specified substitution mutations at positions 16 or 17 in the chymotrypsin numbering system that behave as zymogens until incorporated into prothrombinase complex.

An FXa variant of the disclosure may be produced by any technique for expressing a protein.

An "isolated protein," "isolated polypeptide" or "isolated variant" is a protein, polypeptide or variant that by virtue of its origin or source of derivation (1) is not associated with naturally associated components that accompany it in its native state, (2) is free of other proteins from the same species, (3) is expressed by a cell from a different species, or (4) does not occur in nature. Thus, a polypeptide that is chemically synthesized or synthesized in a cellular system different from the cell from which it naturally originates will be "isolated" from its naturally associated components. A protein may also be rendered substantially free of naturally-

associated components by isolation, using protein purification techniques well known in the art.

A protein or polypeptide is “substantially pure,” “substantially homogeneous,” or “substantially purified” when at least about 60 to 75% of a sample exhibits a single species of polypeptide. The polypeptide or protein may be monomeric or multimeric. A substantially pure polypeptide or protein will typically comprise about 50%, 60%, 70%, 80% or 90% W/W of a protein sample, more usually about 95%, and may be over 99% pure. Protein purity or homogeneity may be indicated by a number of means well known in the art, such as polyacrylamide gel electrophoresis of a protein sample, followed by visualizing a single polypeptide band upon staining the gel with a stain well known in the art. For certain purposes, higher resolution may be provided by using HPLC or other means well known in the art for purification.

The methods of the disclosure are useful to counteract a direct FXa inhibitor. A direct FXa inhibitor is an inhibitor that binds directly to FXa and selectively binds FXa over other proteases. Direct FXa inhibitors are noncompetitive inhibitors of FXa with respect to prothrombin. They bind the substrate binding cleft and inhibit FXa competitively with respect to small peptide substrates that also bind this region. They inhibit FXa with high picomolar affinity and are highly protein bound in plasma. Examples of direct FXa inhibitors are rivaroxaban, apixaban, betrixaban, darexaban, edoxaban and otamixaban. In certain embodiments, direct FXa inhibitors are selected from rivaroxaban and apixaban.

According to the disclosure, an FXa variant can be used to counteract a direct FXa inhibitor that binds FXa or that binds FXa that has formed prothrombinase. The direct FXa inhibitors may or may not require cofactors of FXa for inhibition. According to the methods of the disclosure, an FXa variant, such as FXa^{T16L} and FXa^{T16Z}, are administered to a subject whose blood contains a direct FXa inhibitor.

The disclosure encompasses the use of a FXa variant to counteract direct FXa inhibitors, including but not limited to synthetic inhibitors, small molecule inhibitors, orally available inhibitors, or reversible inhibitors. The FXa inhibitor may be any combination of these features, such as an orally available, synthetic, reversible, small molecule inhibitor. In certain embodiments, the direct FXa inhibitors may be selected from rivaroxaban, apixaban, betrixaban, darexaban, edoxaban and otamixaban (see Perzborn et al., *Nat Rev Drug Discov.* 2011 January; 10(1):61-75; Turpie, *Arterioscler Thromb Vasc Biol.* 2007 June; 27(6):1238-47; Pinto et al., *Expert Opin. Ther. Patents* 22:645-661 (2012); Pinto, et al., *J. Med. Chem.* 50:5339-5356 (2007), each of which is incorporated by reference herein). In certain embodiments, direct FXa inhibitors are selected from rivaroxaban or apixaban.

In some embodiments, a FXa variant of the disclosure can be administered to a subject to reverse the effects of a direct FXa inhibitor where such inhibitor occurs at therapeutic concentrations. In other embodiments, a FXa variant of the disclosure can be administered to a subject to reverse the effects of a direct FXa inhibitor where such inhibitor occurs at supratherapeutic concentrations. A supratherapeutic concentration is one that is higher than that ordinarily considered required to safely achieve anti-coagulation in a particular subject or class of subjects. Supratherapeutic concentrations of a direct FXa inhibitor can result from accidental or intentional overdose. Supratherapeutic concentrations of a direct FXa inhibitor can also result from unexpected effects in particular subjects, such as an unexpectedly high sensitivity to these drugs, or unexpectedly

slow rate of clearance, due for example to drug interactions or other factors. Determination of what would be a therapeutic concentration or supratherapeutic concentration of direct FXa inhibitor in a particular subject or class of subjects is within the knowledge of those ordinarily skilled in the art.

According to the disclosure, an FXa variant is used to counteract a direct FXa inhibitor or inhibitors that selectively bind FXa over other trypsin-like proteases by at least 5-fold, at least 6-fold, at least 7-fold, at least 10-fold, at least 15-fold, at least 20-fold, at least 25-fold, at least 30-fold, at least 50-fold, at least 100-fold, at least, 500-fold, at least 1,000-fold, at least 5,000-fold or at least 10,000-fold.

The direct FXa inhibitor may bind an FXa variant with a K_i of about 2×10^{-7} M or less. “ K_i ” refers to the inhibitor constant of a particular inhibitor-target interaction, which is the concentration required to produce half maximum inhibition. One can determine the K_i by using methods known in the art. The disclosure contemplates, thus, counteracting a direct FXa inhibitor that binds an FXa variant free of the prothrombinase complex with a K_i of about 2×10^{-8} M or less, about 1×10^{-8} M or less, about 9×10^{-9} M or less, about 8×10^{-9} M or less, about 7×10^{-9} M or less, about 6×10^{-9} M or less, about 5×10^{-9} M or less, about 4×10^{-9} M or less, about 3×10^{-9} M or less, about 2×10^{-9} M or less, about 1×10^{-9} M or less, about 9×10^{-10} M or less, about 8×10^{-10} M or less, about 7×10^{-10} M or less, about 6×10^{-10} M or less, about 5×10^{-10} M or less, about 4×10^{-10} M or less, about 3×10^{-10} M or less, about 2×10^{-10} M or less, about 1×10^{-10} M or less, about 9×10^{-11} M or less, about 8×10^{-11} M or less, about 7×10^{-11} M or less, about 6×10^{-11} M or less, about 5×10^{-11} M or less, about 4×10^{-11} M or less, about 3×10^{-11} M or less, about 2×10^{-11} M or less, about 1×10^{-11} M or less, about 9×10^{-12} M or less, about 8×10^{-12} M or less, about 7×10^{-12} M or less, about 6×10^{-12} M or less, about 5×10^{-12} M or less, about 4×10^{-12} M or less, about 3×10^{-12} M or less, about 0.2×10^{-12} M or less, or about 1×10^{-12} M or less, or less. The direct FXa inhibitor to be counteracted by an FXa variant according to the methods of the disclosure may bind a wild-type FXa with a K_i at least 1.5 fold, at least 2-fold, at least 3-fold, at least 4-fold, at least 5-fold, at least 6-fold, at least 7-fold, at least 10-fold, at least 15-fold, at least 20-fold, at least 25-fold, at least 30-fold, or at least 50-fold less than it binds the FXa variant. The direct FXa inhibitor may bind a wild-type FXa with a K_i of at least 20%, at least 30%, at least 40%, at least 50%, at least 60%, at least 70%, at least 80%, at least 90%, at least 95%, or at (east 99% less than the K_i with an FXa variant free of the prothrombinase complex. The direct FXa inhibitor may bind a prothrombinase complex comprising a wild-type FXa with about the same K_i as it binds a prothrombinase complex comprising an FXa variant.

In one aspect, the disclosure provides methods for counteracting the effects of a direct FXa inhibitor in a subject who is bleeding (internally or externally) or is at risk of bleeding (e.g., in the course of a planned surgery) by administering a FXa variant. In some embodiments, the direct FXa inhibitor may be present in the subject at a therapeutic concentration or a higher concentrations (i.e., a supratherapeutic concentration). In some embodiments, the therapeutic concentration may be an overdose in sensitive individuals. The methods of the disclosure, thus, are useful for providing an antidote to an overdose of a direct FXa inhibitor. In various embodiments, the subject of treatment may be a human or a veterinary subject.

Direct inhibitor overdose can be detected based on existence of symptoms or signs of excessively reduced clotting

ability. Non-limiting examples include evidence of gastrointestinal bleeding, including dark tarry stools, bloody stools, and vomiting of blood. Other examples include nosebleeds, and increased tendency to, or severity of, bruising or bleeding from minor cuts and scrapes.

In a clinical setting, direct inhibitor overdose can be detected directly or by measuring the ability of subject blood to clot and detecting deviations from the expected degree of anti-coagulation. Blood clotting potential can be measured in ways familiar to those ordinarily skilled in the art. For example, overdose may be suspected when a subject's prothrombin time is excessively prolonged. In certain embodiments, overdose is confirmed when the prothrombin time expressed as an International Normalized Ratio (INR) is measured to be greater than about 1.0, 1.5, 2.0, 2.5, 3.0, 3.5, 4.0, 4.5, 5.0, 5.5, 6.0, 6.5, 7.0, 7.5, 8.0, 8.5, 9.0, 9.5, 10, 12, 14, 16, 18, 20, or greater.

The FXa variant may be administered whenever it is desired to counteract the effects of the direct FXa inhibitor, including but not limited to before a planned surgery, after an injury resulting in external or internal bleeding or after a direct FXa inhibitor overdose. According to the disclosure, the FXa variant may be administered at least about 12 hours, at least about 6 hours, at least about 3 hours, at least about 2 hours, at least about 1 hour, at least about 30 minutes, at least about 10 minutes, or at least about 5 minutes of when the desired counteracting effect is needed, such as before a planned surgery, after an injury resulting in external or internal bleeding or after a direct FXa inhibitor overdose.

According to another embodiment, the disclosure provides a method of administering a FXa variant to effect the urgent reversal of acquired coagulopathy due to FXa inhibition therapy in a subject with acute major bleeding. In some embodiments, subjects are adult human patients: In other embodiments, subjects are pediatric human patients.

In some embodiments, acute major bleeding is caused by trauma. In other embodiments, acute major bleeding occurs during surgery or other type of interventional procedure. Exemplary non-limiting interventional procedures include incisions, drainage, vascular surgery, appendectomy, herniotomy or hernioplasty, abdominal surgery, cholecystectomy, trephination (burr hole), lumbar puncture, cardiac pacemaker insertion, hip fracture surgery, and others. In yet other embodiments, acute major bleeding can be spontaneous bleeding with no apparent cause.

Without limitation, sites of acute major bleeding include gastrointestinal bleeding, subcutaneous or intramuscular bleeding, bladder bleeding, hemarthrosis, subdural hematoma, nasal bleeding, peritoneal bleeding, uterine bleeding, and other sites of bleeding.

Effective treatment with FXa variants of the disclosure can reverse the effects of a direct FXa inhibitor. Successful reversal of such effects by a FXa variant can be determined in a variety of ways and be measured or monitored using different assays, methods, or endpoints.

In some embodiments, treatment with a FXa variant to reverse the effects of a direct FXa inhibitor is monitored using tests or assays performed on blood or plasma from a subject treated with FXa variant. A blood sample can be taken from a subject at a predetermined time after treatment with FXa variant. The blood, or plasma prepared from it, is then subjected to one or more tests to determine if certain hemostatic pharmacodynamic parameters have been normalized despite the presence of direct FXa inhibitor. If normalization is found then the subject need not be further treated with FXa variant. If normalization is not found, however, then further treatment with FXa variant in accor-

dance with the methods of the disclosure may be required to reverse the effect of a direct FXa inhibitor. Tests for monitoring the effectiveness of treatment with a FXa variant include tests that directly or indirectly measure the ability to clot or that measure the activity of a direct FXa inhibitor. Non-limiting exemplary tests include prothrombin time or the related International Normalized Ratio, the prothrombinase-induced clotting time assay, thromboelastometry, thromboelastography, chromogenic anti-FXa assay, thrombin generation assay, level of prothrombin fragment 1+2, level of thrombin-antithrombin III complex, activated partial thromboplastin time, and partial thromboplastin time. Other tests are also possible within the knowledge of those of ordinary skill in the art.

According to some embodiments, reversing the effects of a direct FXa inhibitor in a subject by administering a FXa variant reduces bleeding in the subject. In some embodiments, treatment with FXa variant reduces bleeding in a subject at least 10%, 20%, 30%, 40%, 50%, 60%, 70%, 80%, 90%, 95%, or 99% in the presence of a direct FXa inhibitor compared to absence of treatment with FXa variant. In other embodiments, treatment with FXa variant reduces bleeding in a subject about 5%-10%, 10%-15%, 15%-20%, 20%-25%, 25%-30%, 30%-35%, 35%-40%, 40%-45%, 45%-50%, 50%-55%, 55%-60%, 60%-65%, 65%-70%, 70%-75%, 75%-80%, 80%-85%, 85%-90%, 90%-95%, or 95%-100%.

According to some embodiments, reversing the effects of a direct FXa inhibitor in a subject by administering a FXa variant reduces the activity of a direct FXa inhibitor in the subject. In some embodiments, treatment with FXa variant reduces activity of the direct FXa inhibitor in a subject at least 10%, 20%, 30%, 40%, 50%, 60%, 70%, 80%, 90%, 95%, or 99% in the presence of a direct FXa inhibitor compared to absence of treatment with FXa variant. In other embodiments, treatment with FXa variant reduces the activity of a direct FXa inhibitor in a subject about 5%-10%, 10%-15%, 15%-20%, 20%-25%, 25%-30%, 30%-35%, 35%-40%, 40%-45%, 45%-50%, 50%-55%, 55%-60%, 60%-65%, 65%-70%, 70%-75%, 75%-80%, 80%-85%, 85%-90%, 90%-95%, or 95%-100%.

Activity of a direct FXa inhibitor can be monitored using a chromogenic anti-FXa assay, such as that described in Asmis, et al., *Thromb Res.*, 129:492-498 (2012), or Barrett, et al., *Thromb Haemost.* 104:1263-71 (2010), each of which are incorporated by reference herein.

According to some embodiments, reversing the effects of a direct FXa inhibitor in a subject by administering a FXa variant increases the amount of thrombin produced in the blood or plasma of the subject. In some embodiments, treatment with FXa variant increases thrombin production in a subject at least 10%, 20%, 30%, 40%, 50%, 60%, 70%, 80%, 90%, 95%, 100%, 1.5 fold, 2-fold, 3-fold, 4-fold, 5-fold, 6-fold, 7-fold, 10-fold, 15-fold, 20-fold, 25-fold, 30-fold, at least 50-fold, or more in the presence of a direct FXa inhibitor compared to the absence of an FXa variant. Thrombin production in the blood or plasma of a subject can be determined using the thrombin generation assay (TGA) or other technique familiar to those of ordinary skill in the art.

According to some embodiments, reversing the effects of a direct FXa inhibitor in a subject by administering a FXa variant increases clotting in the subject. In some embodiments, treatment with FXa variant increases clotting in a subject at least 10%, 20%, 30%, 40%, 50%, 60%, 70%, 80%, 90%, 95%, 100%, 1.5 fold, 2-fold, 3-fold, 4-fold, 5-fold, 6-fold, 7-fold, 10-fold, 15-fold, 20-fold, 25-fold,

30-fold, at least 50-fold, or more in the presence of a direct FXa inhibitor compared to the absence of an FXa variant.

According to some embodiments, reversing the effects of a direct FXa inhibitor in a subject by administering a FXa variant reduces clotting time in the subject. In some embodiments, treatment with FXa variant reduces clotting time in a subject at least 10%, 20%, 30%, 40%, 50%, 60%, 70%, 80%, 90%, 95%, or 99% in the presence of a direct FXa inhibitor compared to absence of treatment with FXa variant. In other embodiments, treatment with FXa variant reduces clotting time in a subject about 5%-10%, 10%-15%, 15%-20%, 20%-25%, 25%-30%, 30%-35%, 35%-40%, 40%-45%, 45%-50%, 50%-55%, 55%-60%, 60%-65%, 65%-70%, 70%-75%, 75%-80%, 80%-85%, 85%-90%, 90%-95%, or 95%-100%.

According to some embodiments, clotting time is determined by measuring the subject's prothrombin time (PT) which decreases as hemostasis is restored. PT is the amount of time it takes for serum to clot after addition of tissue factor. PT therefore measures the capability of the extrinsic clotting system to support clotting. PT can vary depending on the particular reagents a lab uses to run the test, but a normal PT is about 11 to 13 seconds. Clotting time can also be expressed using the International Normalized Ratio (INR), which eliminates lab to lab variability in clotting time measurements. Using the INR, a ratio of 0.8 to 1.1 indicates normal clotting. PT or INR can be determined at a predetermined time after a FXa variant is administered to a subject in need of reversal of the effects of a direct FXa inhibitor.

In some embodiments, treatment with a FXa variant to reverse the effects of a direct FXa inhibitor reduces the PT of a subject to about 25 seconds, 24 seconds, 23 seconds, 22 seconds, 21 seconds, 20 seconds, 19 seconds, 18 seconds, 17 seconds, 16 seconds, 15 seconds, 14 seconds, 13 seconds, 12 seconds, 11 seconds, 10 seconds, or less. In other embodiments, treatment with a FXa variant reduces the INR of a subject to about 4.0, 3.9, 3.8, 3.7, 3.6, 3.5, 3.4, 3.3, 3.2, 3.1, 3.0, 2.9, 2.8, 2.7, 2.6, 2.5, 2.4, 2.3, 2.2, 2.1, 2.0, 1.9, 1.8, 1.7, 1.6, 1.5, 1.4, 1.3, 1.2, 1.1, 1.0, 0.9, 0.8, 0.7, or less. According to other embodiments, treatment with FXa variant reduces PT or INR in a subject about 5%-10%, 10%-15%, 15%-20%, 20%-25%, 25%-30%, 30%-35%, 35%-40%, 40%-45%, 45%-50%, 50%-55%, 55%-60%, 60%-65%, 65%-70%, 70%-75%, 75%-80%, 80%-85%, 85%-90%, 90%-95%, or 95%-100%.

Prothrombin time can be measured at a predetermined after administration of a FXa variant. Thus, in some non-limiting embodiments, PT is measured 15 mins, 20 mins, 30 mins, 40 mins, 45 mins, 50 mins, 60 mins or more after administration of FXa. Other times are also possible according to the knowledge of those of ordinary skill in the art.

Clotting time can also be measured using the One-step prothrombinase-induced clotting time (PiCT) assay as described in Graff, et al., Monitoring effects of direct FXa-inhibitors with a new one-step prothrombinase-induced clotting time (PiCT) assay: comparative in vitro investigation with heparin, enoxaparin, fondaparinux and DX 9065a, *Int J Clin Pharmacol Ther.*, 45:237-43 (2007) and Harder, et al., Monitoring direct FXa-inhibitors and fondaparinux by Prothrombinase-induced. Clotting Time (PiCT): relation to FXa-activity and influence of assay modifications, *Thromb Res.*, 123:396-403 (2008), each of which are incorporated by reference.

In yet other embodiments, the methods of thromboelastometry or thromboelastography may be used to analyze clot formation or clotting time.

According to some embodiments, reversing the effects of a direct FXa inhibitor in a subject by administering a FXa variant increases the level of prothrombin fragment 1+2 (PF1+2) in the blood or plasma of the subject. In some embodiments, treatment with FXa variant increases PF1+2 in a subject at least 10%, 20%, 30%, 40%, 50%, 60%, 70%, 80%, 90%, 95%, 100%, 1.5-fold, 2-fold, 3-fold, 4-fold, 5-fold, 6-fold, 7-fold, 10-fold, 15-fold, 20-fold, 25-fold, 30-fold, at least 50-fold, or more in the presence of a direct FXa inhibitor compared to the absence of an FXa variant.

According to some embodiments, reversing the effects of a direct FXa inhibitor in a subject by administering a FXa variant increases the level of thrombin-antithrombin III complex (TAT) in the blood or plasma of the subject. In some embodiments, treatment with FXa variant increases TAT in a subject at least 10%, 20%, 30%, 40%, 50%, 60%, 70%, 80%, 90%, 95%, 100%, 1.5-fold, 2-fold, 3-fold, 4-fold, 5-fold, 6-fold, 7-fold, 10-fold, 15-fold, 20-fold, 25-fold, 30-fold, at least 50-fold, or more in the presence of a direct FXa inhibitor compared to the absence of an FXa variant.

According to some embodiments, reversing the effects of a direct FXa inhibitor in a subject by administering a FXa variant reduces activated partial thromboplastin time (aPTT) in the subject. In some embodiments, treatment with FXa variant reduces activated partial thromboplastin time (aPTT) in a subject at least 10%, 20%, 30%, 40%, 50%, 60%, 70%, 80%, 90%, 95%, or 99% in the presence of a direct FXa inhibitor compared to absence of treatment with FXa variant. In other embodiments, treatment with FXa variant reduces aPTT in a subject about 5%-10%, 10%-15%, 15%-20%, 20%-25%, 25%-30%, 30%-35%, 35%-40%, 40%-45%, 45%-50%, 50%-55%, 55%-60%, 60%-65%, 65%-70%, 70%-75%, 75%-80%, 80%-85%, 85%-90%, 90%-95%, or 95%-100%.

According to some embodiments, reversing the effects of a direct FXa inhibitor in a subject by administering a FXa variant reduces partial thromboplastin time (PTT) in the subject. In some embodiments, treatment with FXa variant reduces partial thromboplastin time (PTT) in a subject at least 10%, 20%, 30%, 40%, 50%, 60%, 70%, 80%, 90%, 95%, or 99% in the presence of a direct FXa inhibitor compared to absence of treatment with FXa variant. In other embodiments, treatment with FXa variant reduces PTT in a subject about 5%-10%, 10%-15%, 15%-20%, 20%-25%, 25%-30%, 30%-35%, 35%-40%, 40%-45%, 45%-50%, 50%-55%, 55%-60%, 60%-65%, 65%-70%, 70%-75%, 75%-80%, 80%-85%, 85%-90%, 90%-95%, or 95%-100%.

In other embodiments, clinical endpoints can be relied upon to determine if hemostasis has been adequately restored in a subject treated with a FXa variant to reverse the effects of a direct FXa inhibitor. For example, where a subject presents with acute bleeding, clinical hemostatic efficacy can be scored "very good" where prompt cessation of existing bleeding occurs after treatment with FXa variant; "satisfactory" where there is a 1-2 hr delay in bleeding cessation; "questionable" where there is a >2 hr delay in bleeding cessation; and "none" where an effect on bleeding is absent. Where treatment with FXa variant is determined to be less than satisfactory, then an additional dose of FXa variant can be administered to effect adequate hemostasis. In a further example, where a subject is undergoing an interventional procedure, clinical hemostatic efficacy can be scored "very good" where normal hemostasis is attained during the procedure; "satisfactory" where intraprocedural hemostasis is mildly abnormal as judged by quantity or quality of blood loss (e.g., slight oozing); "questionable"

where intraprocedural hemostasis is moderately abnormal as judged by quantity or quality of blood loss (e.g., controllable bleeding); and “none” where intraprocedural hemostasis is severely abnormal as judged by quantity or quality of blood loss (e.g., severe refractory hemorrhage).

A therapeutically effective dose of a direct FXa inhibitor depends upon numerous factors that are well known to a medical practitioner of skill in the art. A typical therapeutic plasma concentration of rivaroxaban is about 500 nM. However, according to the disclosure, an FXa variant can be administered to counteract lower or higher concentrations of inhibitor. The plasma concentration of rivaroxaban in a subject to be treated with an FXa variant may be lower or higher than the typical therapeutic concentration, for example about 100 nM, about 200 nM, about 300 nM, about 400 nM, about 500 nM, about 600 nM, about 700 nM, about 800 nM, about 900 nM or about 1,000 nM.

A typical therapeutic plasma concentration of apixaban is about 250 nM. In certain embodiments, the FXa variant is administered to a subject with a plasma concentration of apixaban of about 100 nM, about 200 nM, about 300 nM, about 400 nM, about 500 nM, about 600 nM, about 700 nM, about 800 nM, about 900 nM or about 1,000 nM.

Likewise, according to the disclosure, an FXa variant can be used to counteract a direct FXa inhibitor in cases of overdose, such as when the plasma concentration of the inhibitor is at least 20%, at least 30%, at least 40%, at least 50%, at least 60%, at least 70%, at least 80%, at least 90%, at least 95%, at least 99%, or at least 1.5 fold, at least 2-fold, at least 3-fold, at least 4-fold, at least 5-fold, at least 6-fold, at least 7-fold, at least 10-fold, at least 15-fold, at least 20-fold, at least 25-fold, at least 30-fold, or at least 50-fold higher than the typical therapeutic plasma concentration.

The FXa variants are surprisingly effective in counteracting a direct FXa inhibitor at a plasma concentration that is lower than the plasma concentration of the direct FXa inhibitor. According to the disclosure, the FXa variant counters the effect of a direct FXa inhibitor at a plasma concentration ratio of variant to inhibitor of about 1 to 10, about 1 to 25, about 1 to 50, about 1 to 100, about 1 to 250, about 1 to 500, about 1 to 1,000, about 1 to 2,500, about 1 to 5,000 or about 1 to 10,000. In certain embodiments, the FXa variant counters the effect of a direct FXa inhibitor at a plasma concentration of at least 10-fold, at least 25-fold, at least 50-fold, at least 100-fold, at least 250-fold, at least 500-fold, at least 1,000-fold, at least 2,500-fold, at least 5,000-fold, or at least 10,000-fold lower than the plasma concentration of the direct FXa inhibitor.

In other embodiments, the plasma concentration of an FXa variant sufficient to reverse the effect of a direct FXa inhibitor is calculated by multiplying the plasma concentration of the direct inhibitor by a conversion factor ranging from about 0.1×10^{-4} to about 1000×10^{-4} , about 4×10^{-4} to about 40×10^{-4} , about 20×10^{-4} to about 200×10^{-4} , or other ranges. In yet other embodiments, the conversion factor can be about 0.1×10^{-4} , 0.5×10^{-4} , 1×10^{-4} , 2×10^{-4} , 3×10^{-4} , 4×10^{-4} , 5×10^{-4} , 6×10^{-4} , 7×10^{-4} , 8×10^{-4} , 9×10^{-4} , 10×10^{-4} , 11×10^{-4} , 12×10^{-4} , 13×10^{-4} , 14×10^{-4} , 15×10^{-4} , 16×10^{-4} , 17×10^{-4} , 18×10^{-4} , 19×10^{-4} , 20×10^{-4} , 21×10^{-4} , 22×10^{-4} , 23×10^{-4} , 24×10^{-4} , 25×10^{-4} , 26×10^{-4} , 27×10^{-4} , 28×10^{-4} , 29×10^{-4} , 30×10^{-4} , 31×10^{-4} , 32×10^{-4} , 33×10^{-4} , 34×10^{-4} , 35×10^{-4} , 36×10^{-4} , 37×10^{-4} , 38×10^{-4} , 39×10^{-4} , 40×10^{-4} , 45×10^{-4} , 50×10^{-4} , 55×10^{-4} , 60×10^{-4} , 65×10^{-4} , 70×10^{-4} , 75×10^{-4} , 80×10^{-4} , 85×10^{-4} , 90×10^{-4} , 95×10^{-4} , 100×10^{-4} , 110×10^{-4} , 120×10^{-4} , 130×10^{-4} , 140×10^{-4} , 150×10^{-4} , 160×10^{-4} , 170×10^{-4} , 180×10^{-4} , 190×10^{-4} , 200×10^{-4} , 250×10^{-4} , 300×10^{-4} , 350×10^{-4} , 400×10^{-4} , 450×10^{-4} , 500×10^{-4} , $550 \times$

10^{-1} , 600×10^{-4} , 650×10^{-4} , 700×10^{-4} , 750×10^{-4} , 800×10^{-4} , 850×10^{-4} , 900×10^{-1} , 950×10^{-4} , or 1000×10^{-4} , and ranges among these numbers. Plasma concentration of FXa direct inhibitor can be measured according to the knowledge of the skilled artisan, for example, by radio-immuno assay (RIA) or other method.

Achieving a target plasma concentration of FXa variant sufficient to reverse overdose of a direct FXa inhibitor is within the knowledge of those ordinarily skilled in the art. In a non-limiting example, estimates of relevant pharmacokinetic parameters, such as subject plasma volume or other parameters, can be made based on upon subject sex, height and weight, or other factors, and used to calculate how much FXa variant needs be administered to achieve the target concentration. After administering FXa variant, plasma concentrations can be monitored according to the knowledge of those ordinarily skilled in the art and this information used to maintain the concentration in any desired range.

The compositions and methods of the disclosure include a “therapeutically effective amount” or a “prophylactically effective amount” of an FXa variant. A “therapeutically effective amount” refers to an amount effective, at dosages and for periods of time necessary, to achieve the desired therapeutic result. A therapeutically effective amount of the FXa variant may vary according to factors such as the disease state, age, sex, and weight of the individual, and the ability of the FXa variant to elicit a desired response in the individual. A therapeutically effective amount is also one in which any toxic or detrimental effects of the FXa variant are outweighed by the therapeutically beneficial effects. A “prophylactically effective amount” refers to an amount effective, at dosages and for periods of time necessary, to achieve the desired prophylactic result. For example, a dose may be given prior to a planned surgery.

Dosage regimens can be adjusted to provide the optimum desired response (e.g., a therapeutic or prophylactic response). For example, a single bolus can be administered, several divided doses can be administered over time or the dose can be proportionally reduced or increased as indicated by the exigencies of the therapeutic situation. It is especially advantageous to formulate parenteral compositions in dosage unit form for ease of administration and uniformity of dosage. Dosage unit form as used herein refers to physically discrete units suited as unitary dosages for the mammalian subjects to be treated; each unit containing a predetermined quantity of active compound calculated to produce the desired therapeutic effect in association with the required pharmaceutical carrier. The specification for the dosage unit forms of the disclosure are dictated by and directly dependent on (a) the unique characteristics of the FXa variant and the particular therapeutic or prophylactic effect to be achieved, and (b) the limitations inherent in the art of compounding such an FXa variant for the treatment of individuals.

In certain embodiments, a therapeutically or prophylactically-effective amount of an FXa variant administered is about 0.0001 to 50 mg/kg, about 0.001 to 50 mg/kg, about 0.001 to 5 mg/kg, about 0.001 to 0.5 mg/kg, about 0.001 to 0.05 mg/kg, about 0.01 to 5 mg/kg or about 0.01 to 0.5 mg/kg.

In certain embodiments, a therapeutically or prophylactically-effective serum concentration of an FXa variant of the disclosure is about 0.0003 to 300 nM, about 0.003 to 300 nM, about 0.03 to 300 nM, about 0.003 to 30 nM, about 0.03 to 30 nM or about 0.3 to 3 nM. The concentration of the

FXa variant, for example in blood or plasma, may be measured by any method known in the art.

It is to be noted that dosage values may vary with FXa inhibitor concentration. It is to be further understood that for any particular subject, specific dosage regimens should be adjusted over time according to the individual need and the professional judgment of the person administering or supervising the administration of the compositions, and that dosage ranges set forth herein are exemplary only and are not intended to limit the scope or practice of the claimed composition.

Another aspect of the present disclosure provides kits comprising an FXa variant or a composition comprising such an FXa variant. A kit may include, in addition to the FXa variant or composition, diagnostic or additional therapeutic agents. A kit can also include instructions for use in a therapeutic method, as well as packaging material such as, but not limited to, ice, dry ice, styrofoam, foam, plastic, cellophane, shrink wrap, bubble wrap, cardboard and starch peanuts. In one embodiment, the kit includes the FXa variant or a composition comprising it and one or more therapeutic agents that can be used in a method described herein.

The FXa variant may be administered, for example in a composition comprising it, once or multiple times to a subject until adequate hemostasis is restored or the direct FXa inhibitor or inhibitors are no longer effective. Where multiple administrations are used they may be administered hourly, daily, or at any other appropriate interval, including for example multiple daily doses. Multiple doses may be administered on a schedule such as every 10 minutes, every 15 minutes, every 20 minutes, every 30 minutes, every hour, every two hours, every three hours, every four hours, three times daily, twice daily, once daily, once every two days, once every three days, and once weekly. The FXa variant may also be administered continuously, e.g. via a minipump. The FXa variant may be administered, for example, via a parenteral route (e.g., intravenously, subcutaneously, intraperitoneally, or intramuscularly). The FXa variant will generally be administered as part of a pharmaceutical composition as described below.

In another embodiment, the FXa variant may be co-administered with another procoagulant including another FXa variant, Factor IX, Factor XIa, Factor XIIa, Factor VIII, Factor Vila, FEIBA and prothrombin complex concentrate (PCC).

Co-administration of an FXa variant of the disclosure with an additional therapeutic agent (combination therapy) encompasses administering a pharmaceutical composition comprising the FXa variant and the additional therapeutic agent, as well as administering two or more separate pharmaceutical compositions, i.e., one comprising the FXa variant and the other(s) comprising the additional therapeutic agent(s). Co-administration or combination therapy further includes administering the FXa variant and additional therapeutic agent(s) simultaneously or sequentially, or both. For instance, the FXa variant may be administered once every three days, while the additional therapeutic agent is administered once daily at the same as the FXa variant, or at a different time. An FXa variant may be administered prior to or subsequent to treatment with the additional therapeutic agent. Similarly, administration of an FXa variant of the disclosure may be part of a treatment regimen that includes other treatment modalities including surgery. The combination therapy may be administered to prevent recurrence of the condition. The combination therapy may be administered from multiple times hourly to weekly. The administrations may be on a schedule such as every 10 minutes,

every 15 minutes, every 20 minutes, every 30 minutes, every hour, every two hours, every three hours, every four hours, three times daily, twice daily, once daily, once every two days, once every three days, once weekly, or may be administered continuously, e.g. via a minipump. The combination therapy may be administered, for example, via a parenteral route (e.g., intravenously, subcutaneously, intraperitoneally, or intramuscularly).

In a further aspect, the disclosure provides a composition comprising an FXa variant for use in counteracting a direct FXa inhibitor in a subject. The composition may comprise a pharmaceutically acceptable carrier, vehicle or other ingredients that are physiologically compatible. Non-limiting examples of such carriers, vehicles and other ingredients include solvents (e.g., water, ethanol, saline, phosphate buffered saline), detergents, surfactants, dispersion media, coatings, antibacterial or antifungal agents, isotonic agents, absorption delaying agents, sugars (e.g., sucrose, dextrose, lactose), polyalcohols (e.g., glycerol, mannitol, sorbitol), salts (e.g., sodium chloride, potassium chloride), wetting agents, emulsifying agents, preservatives, buffers, and agents capable of enhancing the stability or effectiveness of the FXa variant.

A composition for use according to the disclosure may be in any suitable form for administration to a subject, such as liquid solutions (e.g., injectable and infusible solutions). Compositions can be provided in a pre-mixed format ready for administration to a subject, for example, in a vial or pre-filled syringe. Such formats do not require reconstitution with diluent before administration. Alternatively, compositions can be provided in lyophilized form requiring reconstitution with diluent (e.g., sterile water or saline) before administration. If the latter, diluent can be provided with the lyophilisate in a separate container. According to the knowledge of those of ordinary skill in the art, compositions can be formulated for storage under refrigeration or at room temperature. The form of the composition depends, at least in part, on the intended mode of administration. In certain embodiments, the mode of administration is parenteral, including for example intravenous, subcutaneous, intraperitoneal, or intramuscular administration.

Therapeutic compositions typically must be sterile and stable under the conditions of manufacture and storage. The composition can be formulated as a solution, microemulsion, dispersion, in liposomes, or other ordered structure suitable to high drug concentration. Sterile injectable solutions can be prepared by incorporating the FXa variant in the required amount in an appropriate solvent with one or a combination of ingredients enumerated above, as required, followed by filtered sterilization. Generally, dispersions are prepared by incorporating the active compound into a sterile vehicle that contains a basic dispersion medium and the required other ingredients from those enumerated above. In the case of sterile powders for the preparation of sterile injectable solutions, the preferred methods of preparation are vacuum drying and freeze-drying that yields a powder of the active ingredient plus any additional desired ingredient from a previously sterile-filtered solution thereof. The proper fluidity of a solution can be maintained, for example, by the use of a coating such as lecithin, by the maintenance of the required particle size in the case of dispersion and by the use of surfactants. Prolonged absorption of injectable compositions can be brought about by including in the composition an agent that delays absorption, for example, monostearate salts and gelatin.

It is further contemplated by the present disclosure that any of the compositions herein may be administered to a subject being treated with a direct FXa inhibitor.

It is understood that the examples and embodiments described herein are for illustrative purposes only and that various modifications or changes in light thereof will be apparent to persons skilled in the art and are to be included within the and can be made without departing from the true scope of the invention.

EXAMPLES

Example 1—FXa^{T16L} Sensitivity Toward Rivaroxaban

To test the sensitivity of FXa^{T16L} toward rivaroxaban, inhibition assays were established. Rivaroxaban was an efficient inhibitor of wild-type FXa exhibiting an inhibition constant (K_i) of 0.582 nM (FIG. 1A). Due to the zymogen-like nature of FXa^{T16L}, rivaroxaban bound with a ~15-fold reduced affinity to this variant (K_i =9.3 nM) (FIG. 1B). In contrast, when the variant was assembled in the prothrombinase complex (i.e. upon addition of FVa and phospholipid vesicles), the K_i for rivaroxaban was nearly restored to a value comparable to the wild-type enzyme (wt FXa, K_i =2.67 nM (FIG. 2A); FXa^{T16L}, K_i =3.4 nM (FIG. 2B).

Example 2—FXa Variants Counteract Rivaroxaban and Apixaban

The thrombin generation assay (TGA) was used to assess whether the zymogen-like FXa variants can reverse the effects of direct FXa inhibitors in a more physiologic environment. The TGA measures thrombin production in plasma over time following the initiation of coagulation and was performed as previously described (See Bunce et al., (2011) *Blood* 117, 290-298, incorporated by reference herein in their entirety). Thrombin generation in normal human plasma was measured for 90 min at 37° C. in the presence or absence of 500 nM rivaroxaban. To evaluate if FXa^{T16L} can reverse the effect of rivaroxaban, increasing amounts of FXa^{T16L} was added to plasma containing 500 nM rivaroxaban. Thrombin generation was initiated with 2.0 pM tissue factor/4 μ M phospholipid as well as CaCl₂ and a thrombin fluorogenic substrate.

The data demonstrated that the thrombin generation profile of plasma in the presence of 500 nM rivaroxaban was substantially reduced compared to plasma in the absence of rivaroxaban. In contrast, increasing concentrations of FXa^{T16L} from 0.03 to 1 nM restored thrombin generation (FIG. 3). These data show that unexpectedly low concentrations of FXa^{T16L} in the nanomolar and subnanomolar range can reverse the effects of the inhibitor. Dose response analysis of FXa^{T16L} in the presence of 500 nM rivaroxaban (a typical therapeutic plasma concentration) shows that the peak height of thrombin generation (FIGS. 4A and C) and total thrombin produced (ETP) (FIGS. 4 B and D) essentially reached a maximum and was completely restored to normal levels between 1-3 nM of FXa^{T16L} under these conditions. Further experiments showed that even in the presence of high concentration of rivaroxaban (7.5 μ M; supratherapeutic), FXa^{T16L} was still quite effective at a relatively low dose (\leq 3.0 nM) in restoring peak thrombin (FIG. 5A) as well as total thrombin generated (FIG. 5B).

Similar experiments were also performed to evaluate whether FXa zymogen-like variants could reverse the effects of another direct FXa inhibitor, apixaban. In these experi-

ments, the effectiveness of FXa^{T16L} with another zymogen-like FXa variant, FXa^{T16T}, was also compared. FXa^{T16T} is similar to FXa^{T16L}, however it has intrinsically less activity, has a longer plasma half-life, and has ~3-5-fold reduced activity compared to FXa^{T16L} when assembled in the prothrombinase complex. Consistent with the rivaroxaban data, FXa^{T16L} could restore peak thrombin (FIG. 6A) and total thrombin (FIG. 6B) generated in the presence of 250 nM apixaban (a typical therapeutic plasma concentration) in a dose-dependent manner, which appears to reach a maximum between 1-3 nM of FXa^{T16L}. FXa^{T16L} was also effective at reversing the effects of apixaban; however, it appears that higher concentrations of this variant are needed to fully restore thrombin generation (FIG. 6). Both variants were still effective even in the presence of a higher concentration of apixaban (2 μ M). However, under these conditions it appears that higher concentrations of both variants are needed to fully restore thrombin generation (FIGS. 7A and B).

Example 3—FXa^{T16L} Counteracts Rivaroxaban and Apixaban in Whole Blood

Whole blood thromboelastometry was used to assess the ability of the FXa^{T16L} variant to reverse the effects of the direct FXa inhibitors in whole blood. In this system, blood is drawn from healthy volunteers. The first 2 mL of blood were discarded and the subsequent 5 mL of blood was collected into a vacutainer (BID, Franklin Lakes, N.J.). Corn trypsin inhibitor and sodium citrate were in the collection tube, prior to collection of the blood sample, to achieve a final concentration of 0.105 M citrate and 25 μ g/mL corn trypsin inhibitor (Haematologic Technologies, Burlington, Vt.) in the blood. Two sets of reactions were analyzed for each donor. The initial reaction initiated 5 minutes post blood collection. The second reaction initiated 1 hour post initiation of first reaction (1 hour 5 minutes after collection).

Blood was analyzed using Thromboelastometry ROTEM® delta (Tem International GmbH, Munich, Germany). For the reaction: (1) 6 μ L of vehicle, protein, and/or inhibitor were added to the empty cup, (2) 20 μ L of 0.2 M CaCl₂ (final concentration 11.6 mM), and (3) 20 μ L of Innovin (final concentration in reaction 1:10,000; source of tissue factor) were added to the cup. Whole blood collected as described above was added to the reaction (300 μ L and recordings were allowed to proceed for approximately 30-60 minutes. The data collected was analyzed using the manufacturer's software. (Rotem Gamma Software Version 1.1.1).

The ability of FXa^{T16L} to accelerate whole blood clot formation in the presence of rivaroxaban or apixaban was examined using rotational thromboelastometry (ROTEM). Both direct FXa inhibitors alone and at two different concentrations have a substantial effect on whole blood clot formation: at low doses (therapeutic concentrations), whole blood clot formation is partially eliminated (FIG. 8A and FIG. 9A), while at high doses (supratherapeutic concentrations), whole blood clot formation is almost completely eliminated (FIG. 8B and FIG. 9B). The effects of either rivaroxaban or apixaban on whole blood clot formation could be reversed by FXa^{T16L}. In the presence of either 500 nM rivaroxaban or 250 nM apixaban, 0.3 nM FXa^{T16L} could fully or nearly fully restore whole blood coagulation (FIG. 8A and FIG. 9A). When higher concentrations of the direct FXa inhibitors were used (~2 μ M), 0.3 nM FXa^{T16L} partially restored whole blood coagulation and 3 nM FXa^{T16L} fully restored it (FIG. 8B and FIG. 9B). These data demonstrated that an FXa zymogen-like variant can effectively reverse the

anticoagulant effect of rivaroxaban or apixaban in plasma-based and whole-blood coagulation assays at both therapeutic and supratherapeutic concentrations of the inhibitor.

The results of these studies were confirmed and extended by testing if FXa^{T16L} could counteract the anti-coagulant effect of rivaroxaban when both agents were administered in vivo. In these experiments, C57BL/6 mice were infused with rivaroxaban (1 mg/kg) or buffer via the tail vein. Mice were then prepared to expose the jugular vein and the vena cava. Approximately 10 mill later FXa^{T16L} (1 or 2 mg/kg) was infused by direct injection into the jugular vein. Five minutes post injection blood was collected via the vena cava into citrate and corn trypsin inhibitor. Collected blood was then analyzed by ROTEM using dilute tissue factor (Innovin, 1:42,000 dilution). Whole blood from mice administered buffer only clotted by about 2 min (FIG. 12). Administration of 1 mg/kg rivaroxaban substantially prolonged the clot time to about 10 min (FIG. 12). Further administration of FXa^{T16L} shortened clotting time in the presence of rivaroxaban in a dose dependent manner (FIG. 12).

Example 4—FXa^{T16L} Counteracts Rivaroxaban in a Thrombin Generation Assay

The effect of FXa^{T16L} on reversing rivaroxaban in plasma was examined in a thrombin generation assay (TGA) using the calibrated automated thrombography (CAT) system (Thrombinoscope BV, Maastricht, The Netherlands). Normal human plasma was obtained from George King Biomedical (Overland Park, Kans.). In the reaction, 20 μ L of PPP-Reagent LOW containing 4 μ M phospholipids and 1 pM tissue factor was added to 70 μ L of pooled citrated normal human plasma (treated with 250 nM rivaroxaban, within the therapeutic plasma concentration range) in an Immulon 2HB round bottom 0.96 well plate with reactions duplicated. Immediately preceding reaction initiation, 10 μ L of vehicle or FXa^{T16L} was added to plasma at final concentrations ranging from 0.03125 nM to 0.5 nM FXa^{T16L}, given a 120 μ L total reaction volume. Reactions were initiated by addition of 20 μ L FluCa buffer containing calcium chloride and fluorogenic substrate. Fluorescence of plasma reactions was read at 37° C. at 20 second intervals on a Fluoroskan Ascent fluorometer and compared to those of reference thrombin calibrator reactions to determine thrombin concentrations. The intensity of the fluorescence signal (FU) was continuously monitored at 37° C. using the CAT. Thrombin generation curves (nM thrombin vs. time) were analyzed to extract lag time, peak height, time to peak, and the area under the curve representing the endogenous thrombin potential (ETP) using the Thromboscope software (Thrombinoscope BV version).

A dose dependent inhibition of thrombin generation in normal human plasma was observed with in vitro rivaroxaban treatment (5-200 nM) (FIG. 10A). Rivaroxaban resulted in an increase in the lag time coupled with a decrease in the peak thrombin and a decrease in the ETP. The addition of FXa^{T16L} to rivaroxaban (250 nM) inhibited human plasma resulted in a dose dependent reversal of thrombin inhibition (FIG. 10B): peak thrombin generation was restored, the lag phase was shorter, and the ETP increased. At a low dose of 0.03125 nM FXa^{T16L}, thrombin generation was restored to levels comparable to vehicle treated normal human plasma.

Example 5—FXa^{T16L} Counteracts Rivaroxaban in a Mouse Tail Clip Bleeding Model

The ability of FXa^{T16L} to overcome the effects of rivaroxaban in vivo was assessed in an acute bleeding model in

normal mice. The results demonstrated that a zymogen-like FXa variant could reverse the anticoagulant effect of a direct FXa inhibitor.

To establish a dose of rivaroxaban that would prolong bleeding, male C571Bl/6 mice (The Jackson Laboratory, Bar Harbor, Me.) received a single intravenous injection of rivaroxaban at a dose of 10, 25 or 50 mg/kg. Thirty minutes later, mice were anesthetized with isoflurane and placed on a heated platform, and the body temperature of the mice was maintained at 37° C. prior to the tail cut. The tails were immersed in 50 mL pre-warmed phosphate buffered saline (PBS) at 37° C. for 2 minutes. A 3 mm tail cut was made and blood was collected into PBS for a 10 minute period. A quantitative assessment of the amount of bleeding was determined by hemoglobin content of the blood collected into PBS. Tubes were centrifuged to collect erythrocytes, resuspended in 5 mL lysis buffer (8.3 g/L NH₄Cl, 1.0 g/L KHCO₃, and 0.037 g/L EDTA), and the absorbance of the sample was measured at 575 nm. The absorbance values were converted to total blood loss (μ L) using a standard curve. The administration of rivaroxaban resulted in a dose dependent increase in blood loss following a tail cut (FIG. 11).

In this model, a dose of 50 mg/kg rivaroxaban resulted in an increase in blood loss following the tail transection. Mice were dosed with 50 mg/kg rivaroxaban and 30 minutes later 50 or 200 μ g/kg of FXa^{T16L} was dosed intravenously at 37° C. prior to the tail cut. Mice were then anesthetized with isoflurane and placed on a heated platform, and the body temperature of the mice was maintained at 37° C. prior to the tail cut. The tails were immersed in 50 mLs pre-warmed phosphate buffered saline (PBS) at 37° C. for 2 minutes. A 3 mm tail cut was made and blood was collected into PBS for a 10 minute period and the assessment of the amount of bleeding was determined by hemoglobin content as described. In this model, the administration of the hemostatic FXa^{T16L} variant decreased the excessive bleeding loss induced with rivaroxaban (FIG. 11).

Example 6—FXa^{T16L} Counteracts Rivaroxaban in a Mouse Bleeding Model Demonstrated Using Intravital Microscopy

As visualized using intravital microscopy, rivaroxaban was demonstrated to inhibit thrombus formation in the microcirculation of the mouse cremaster muscle after laser-induced injury. Further administration of FXa^{T16L} could counteract the anti-coagulant effect of rivaroxaban in this system.

Using standard techniques, the cremaster muscle of mice was exposed and visualized using intravital microscopy. A vascular injury in the muscle was then induced using a laser. After injury, clot formation was visualized using different fluorescently labeled antibodies that specifically recognize fibrin and platelets. Clotting is indicated by the presence of fluorescent signal from both types of antibodies.

After laser injury, an untreated mouse rapidly formed a clot at the site of injury that was stable for several minutes (FIG. 13A). In the video frame, the clot is visible as the coincidence of fluorescent signal associated with antibodies against fibrin and platelets (light gray center region overlapping darker gray region). Administration of 1 mg/kg rivaroxaban to a mouse, however, delayed the accumulation of platelets at the injury site and eliminated any signs of fibrin (FIG. 13B). In the video frame, only a reduced extent of platelets can be seen as indicated by the dark gray region, which reflects presence of fluorescent signal associated with

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What is claimed is:

1. A method for reversing the anticoagulant effect of a direct Factor Xa inhibitor in a subject, comprising administering to said subject a Factor Xa variant that contains at least one modification selected from the group consisting of:

a) the amino acid at the position corresponding to 235 in SEQ ID NO:1 is substituted with Thr, Leu, Phe, Asp or Gly; and

b) the amino acid at the position corresponding to 236 in SEQ ID NO:1 is substituted with Leu, Ala, or Gly,

wherein said subject has acute major bleeding and acquired coagulopathy due to FXa inhibition therapy using a direct FXa inhibitor,

wherein reversal of the anticoagulant effect of the direct Factor Xa inhibitor effects the urgent reversal of the acquired coagulopathy due to FXa inhibition therapy in said subject with acute major bleeding; and

wherein said Factor Xa variant is effective to reverse the anticoagulant effect of the direct Factor Xa inhibitor at a plasma concentration at least 250-fold lower than the plasma concentration of the direct Factor Xa inhibitor.

2. The method of claim 1, wherein reversal of the anticoagulant effect of the direct Factor Xa inhibitor results in a

45 reduction in bleeding of at least about 5%-10%, 10%-15%, 15%-20%, 20%-25%, 25%-30%, 30%-35%, 35%-40%, 40%-45%, 45%-50%, 50%-55%, 55%-60%, 60%-65%, 65%-70%, 70%-75%, 75%-80%, 80%-85%, 85%-90%, 90%-95%, or 95%-100%.

50 3. The method of claim 1 wherein reversal of the anticoagulant effect of the direct Factor Xa inhibitor results in the amount of thrombin being produced to increase by at least about 10%, 20%, 30%, 40%, 50%, 60%, 70%, 80%, 90%, 95%, 100%, 1.5-fold, 2-fold, 3-fold, 4-fold, 5-fold, 6-fold, 7-fold, 10-fold, 15-fold, 20-fold, 25-fold, 30-fold, or 50-fold.

60 4. The method of claim 1 wherein reversal of the anticoagulant effect of the direct Factor Xa inhibitor results in reduction in clotting time of at least about 5%-10%, 10%-15%, 15%-20%, 20%-25%, 25%-30%, 30%-35%, 35%-40%, 40%-45%, 45%-50%, 50%-55%, 55%-60%, 60%-65%, 65%-70%, 70%-75%, 75%-80%, 80%-85%, 85%-90%, 90%-95%, or 95%-100%.

65 5. The method of claim 4, wherein the reduction in clotting time is measured using prothrombin time (PT).

6. The method of claim 5, wherein said PT in said subject is about 25 seconds, 24 seconds, 23 seconds, 22 seconds, 21

seconds, 20 seconds, 19 seconds, 18 seconds, 17 seconds, 16 seconds, 15 seconds, 14 seconds, 13 seconds, 12 seconds, 11 seconds, or 10 seconds.

7. The method of claim 5, wherein the International Normalized Ratio (INR) in said subject is about 4.0, 3.9, 3.8, 3.7, 3.6, 3.5, 3.4, 3.3, 3.2, 3.1, 3.0, 2.9, 2.8, 2.7, 2.6, 2.5, 2.4, 2.3, 2.2, 2.1, 2.0, 1.9, 1.8, 1.7, 1.6, 1.5, 1.4, 1.3, 1.2, 1.1, 1.0, 0.9, 0.8, or 0.7.

8. The method of claim 5, wherein PT is determined 15 mins, 20 mins, 30 mins, 40 mins, 45 mins, 50 mins, 60 mins, 75 min, or 90 min after administration of the FXa variant.

9. The method of claim 1, wherein the Factor Xa variant is administered before a planned surgery, after an injury or after a direct Factor Xa inhibitor overdose.

10. The method of claim 1, wherein Factor Xa variant is administered more than one time.

11. The method of claim 1, wherein at least one additional procoagulant is administered.

12. The method, composition or use of claim 11, wherein the procoagulant is selected from the group consisting of: a different Factor Xa variant, Factor IX, Factor XIa, Factor XIIa, Factor VIII, Factor VIIa, FEIBA and prothrombin complex concentrate (PCC).

13. The method of claim 1, wherein the plasma concentration of the direct FXa inhibitor is a suprathapeutic amount.

14. The method of claim 1, wherein the direct FXa inhibitor is rivaroxaban, apixaban, betrixaban, darexaban, edoxaban, or otamixaban.

15. The method of claim 1, wherein the plasma concentration of the direct FXa inhibitor is at least about 50 nM, 100 nM, 150 nM, 200 nM, 250 nM, 300 nM, 350 nM, 400 nM, 500 nM, 600 nM, 700 nM, or 800 nM.

16. The method of claim 1, wherein the Factor Xa variant is administered in a dose range of 0.0001 to 50 mg/kg, 0.001 to 50 mg/kg, 0.001 to 5 mg/kg, 0.001 to 0.5 mg/kg, 0.001 to 0.05 mg/kg, 0.01 to 5 mg/kg, or 0.01 to 0.5 mg/kg.

17. The method of claim 1, wherein the Factor Xa variant is administered to achieve a plasma concentration in a range of 0.0003 to 300 nM, 0.003 to 300 nM, 0.03 to 300 nM, 0.003 to 30 nM, 0.03 to 30 nM, or 0.3 to 3 nM.

18. A method for reversing the anticoagulant effect of a direct Factor Xa inhibitor in a subject, comprising administering to said subject a Factor Xa variant comprising Leu at the position corresponding to 235 in SEQ ID NO:1,

wherein said subject has acute major bleeding and acquired coagulopathy due to FXa inhibition therapy using a direct FXa inhibitor,

wherein reversal of the anticoagulant effect of the direct Factor Xa inhibitor effects the urgent reversal of the acquired coagulopathy due to FXa inhibition therapy in said subject with acute major bleeding; and

wherein said Factor Xa variant is effective to reverse the anticoagulant effect of the direct Factor Xa inhibitor at a plasma concentration at least 250-fold lower than the plasma concentration of the direct Factor Xa inhibitor.

19. The method of claim 18, wherein said Factor Xa variant is effective to reverse the anticoagulant effect of the direct Factor Xa inhibitor at a plasma concentration at least 500-fold lower than the plasma concentration of the direct Factor Xa inhibitor.

20. The method of claim 18, wherein said Factor Xa variant is effective to reverse the anticoagulant effect of the direct Factor Xa inhibitor at a plasma concentration at least 1000-fold lower than the plasma concentration of the direct Factor Xa inhibitor.

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